Quest -

Quick User-directed Expression Search Tool

IntelliGenetics

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A;Molecule type: mRNA
A;Residues: 1-139 <HIR>
A;Crose-references: EMBL:X69889; NID:g17733; PID:g17734
A;Experimental source: clone A20
C;Superfamily: beta-1,3-glucanase
S31612 Length: 139 October 13, Found using 'claim36' (zara371.key)
                                                                                                                           R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R. submitted to the EMBL Data Library, December 1992
A;Description: The anther-specific protein encoded by the Brassica napus and Arabidopsis thaliana A6 gene exhibits homology to beta-1,3-glucanases.
A;Reference number: S31612
                                                                                                                                                                                                                           C;Species: Brassica napus (rape)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 17-Nov-2000
                                                                                                                                                                                                                                                                                                         match found in sequence:
831612; TOIG of: 831612
(from "claim36pir.pep")
TOIG of: 831612 check:
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Selected scope is Sequence.
                                                                                                                                                                                                                                                                   F1;S31612 - beta-1,3-glucanase homolog (clone A20) - rape (fragment)
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Time to start comparison
Notify at end of run
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claim36 (λλ) ID claim36 λλ preliminary pattern
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non-matching hits only
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Hit display
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Sequence or key file
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             Type: P
             Check: 219
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BY A; Title: Sequence analysis of the genome of the unicellular cyanobacterium synechocystis sp. PCC6803. II. Sequence determination of the entire genome assignment of potential protein-coding regions.

A; Reference number: S74322; MUID:97061201; PMID:8905231

A; Accession: S74688
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-391 <KAN>
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ound using 'claim36' (zara371.key)
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  of.
                                             of sequences searched: of sequence hits:
                                                                                                                                                                                                                                                    SQMHGGLGLAMLVMSTLLCARYWQGFRVPNKKGAGD
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346 354
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99 107
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separate matches:
sequence hits saved:
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K.; Okumura, S.; Shimpo,
M. Tabata, S.
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aaq23517; LD10322p.
(from "claim36uni.pep")
TOIG of: aaq23617 check: 1097
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02-MAR-2004
02-MAR-2004
02-MAR-2004
              SEQUENCE FROM N.A.

STRAIN-Berkley;

Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J.,

Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J.,

Champe M., Chavez C., Dozsett V., Dreenek D., Farfan D., Friee E.,

George R., Gonzalez M., Guarin H., Kronmiller B., Li P., Liao G.,

Miranda A., Mungall C.J., Nunco J., Pacleb J., Paragas V., Park S.,

Patel S., Phouanenavong S., Wan K., Yu C., Lewis S.E., Rubin G.M.,
                                                                                                             Drosophila melanogaster (Fruit fly).
Bukaryota; Metazoa; Arthropoda; Harapoda; Insecta;
Neoptera; Endopterygota; Diptera; Brachycera; Musco
Ephydroidea; Drosophilidae; Drosophila.
  Submitted
           Patel S., I
Celniker S.
                                                                                                   NCBI_TaxID=7227;
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key from "zara371
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Sequence or key I
List of hits
Hit display
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                             iao G.,
Park S.,
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Yes
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1 match found in sequence:
    cae45695; Hypothetical protein.
    (from "claim36uni.pep")
    TOIG of: cae45695 check: 6871
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aaq72537; SCL-PHA synthase.
(from "claim36uni.pep")
TOIG of: aaq72537 check: 4418
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CAE45695;
02-MAR-2004
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02-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seo S.H., Choi G.G., Rhee Y.H.;
"Cloning of sci-PHA synthase locus and mcl-PHA synthase Pseudomonas sp. HJ-2.";
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY370931; AA072557.1; -.
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AAQ72537;
SEQUENCE FROM N.A. STRAIN=Tu4055;
                                                                                                                                                         Hypothetical protein. Streptomyces parvulus.
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Hypothetical
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Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
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                                                                           NCBI_TaxID=146923;
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MEDLINE=97061201; PubMed=8905231;
MEDLINE=97061201; PubMed=8905231;
Kaneko T., Sato S., Kotani H., Tanaka A., Asamizu E., Nak
Miyajima N., Hirosawa M., Sugiura M., Sasamoto S., Kimura
Miyajima N., Hirosawa M., Sugiura M., Nakazaki N., Naruo K.
Shimpo S., Takeuchi C., Wada T., Watanabe A., Yamada M.,
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                                                                                                                                                                                                                                                     "Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. II. Sequence determination of the centire genome and assignment of potential protein-coding regions."
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Chem. Commun. 22:2780-2788(2003).
EMBL; AJ580915; CAE45695.1; -.
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Synechocystis sp. (strain PCC 6803).
Bacteria; Cyanobacteria; Chroococcales;
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                                                                                                                                                                                                                                                                                                                                                                                      match found in avaitable q6ufw5; SCL-PHA synthase (from "claim36uni.pep")
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              296
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Found usir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P72824
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q06913 ; Beta-1,3-glucanase homologue
(from "claim36uni.pep")
   OCCUPATO DO OCCUPA
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05-JUL-2004
05-JUL-2004
05-JUL-2004
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Q06913;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hird D.L., Worrall D., Hodge R., Smartt S., Paul W., Scott "The anther-specific protein encoded by the Brassica napus Arabidopsis thaliana A6 gene displays similarity to beta-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brassica napus (Rape).

Bukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; eudicotyledons;
eurosids II; Brassicales; Brassicaceae; Brass;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1996 (TrEMBLrel. 01, 0
01-NOV-1996 (TrEMBLrel. 21, 1
01-JUN-2003 (TrEMBLrel. 24, 1
Beta-1,3-glucanase homologue
                                                                                                       SCL-PHA synthase.
Name=phaC;
                                                                                                                                                                                                                                                                                                                            Q6UFW5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plant J
Bacteria; Proteobacteria;
Pseudomonadaceae; Pseudom
                                                                                                                                             SCL-PHA
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                                                                    Pseudomonas sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=94108487; PubMed=8281185;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=3708;
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using 'claim36'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 found in sequence: ; SCL-PHA synthase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FWLPAIAFSWLLGSSILNNGLIPLLLEILQGNQTGVGIGLYFGCVGLAGDVFTRYWSTQS
346 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VWCVAVEGANETELGQALDFACGRSNATCAALAPGRECYAPVSVTWHASYAFSSYWAQFR
107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NQSSQCYFNGLARETTTNPGNEQCKFPSVTL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length: 139 October 13,
ng 'claim36' (zara371.key)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         q06913
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4:1023-1033(1993).
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(zara371.key)
   Pseudomonas
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                              Gammaproteobacteria; Pseudomonadales;
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eudicots; rosid
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P SEQUENCE FROM N.A.

C STRAIN=HJ-2;

A Seo S.H., Choi G.G., Rhee Y.H.;

A Seo S.H., Choi G.G., Rhee Y.H.;

A Seo S.H., Choi G.G., Rhee Y.H.;

L Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.

L Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.

DR InterPro; IPR000073; A/b hydrolase.

DR InterPro; IPR010941; PhaCN.

DR InterPro; IPR010941; PhaCN.

DR Pfam; PP00561; Abhydrolase_1; I.

DR Pfam; PP07167; PhaCN; 1.

DR TIGREAMS; TIGR01838; PHA synth I; 1.

TOR TIGREAMS; TIGR01838; PHA synth I; 1.

CROIENCE 566 AA; 63672 MW; P4CBA7A4CE5943FO CRC64;

CROIENCE 566 AA; 63672 MW; P4CBA7A4CE5943FO CRC64;
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q70hx4; Hypothetical pro
(from "claim36uni.pep")
                                                                                                                                   A LO CONTRACTOR REPORTED TO THE CONTRACTOR REPOR
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Found using
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Found using 'claim36'
                                        Q70HX4 Length: 180
                                                                                                                      InterPro; IPR002048; EF-hand.
InterPro; IPR010983; EF Hand_like.
Pfam; PF00036; efhand; 3.
SMART; SM00054; EFh, 3
PROSITE; PS00018; EF HAND; 2.
Calcium; Calcium-binding; Hypothetic
SEQUENCE 180 AA; 19338 MW; 6CCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q70HX4;
Q70HX4;
05-JUL-2004;
05-JUL-2004;
05-JUL-2004;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Evidence from engineered gene fusions for t
module in a modular polyketide synthase";.";
Chem. Commun. 22:2780-2782 (2003).
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Salas J.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
STRAIN=Tu4055;
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Streptomycineae; Streptomycetaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AJ580915; CAE45695.1;
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q70hx4 check: 6871
    'claim36'
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October 13, 2004 13:25 (zara371.key)
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27,
                                                                                                                               Hypothetical protein.
MW; 6CCE28A1E48436DB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Actinobacteridae; Actinomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from: 1
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Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Streptomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        180
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                                        Type: P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mendez C., Leadlay P.F.,
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                                             Check:
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q7mgu7; Sensor histidir
(from "claim36uni.pep")
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                                                                                                                                                                                    Found
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                                                                                                                                                                                       Q7MGU7 Length: 1156
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GO; GO:0016020; C:membrane; IEA.
GO; GO:0005574; F:ATP binding; IEA.
GO; GO:0000577; F:DNA binding; IEA.
GO; GO:0000577; F:DNA binding; IEA.
GO; GO:00016301; F:kinase activity; IEA.
GO; GO:0000155; F:two-component response regulator activity; IEA.
GO; GO:0000156; F:two-component sensor molecule activity; IEA.
GO; GO:0000155; F:transport; IEA.
GO; GO:0000156; F:two-component sensor molecule activity; IEA.
GO; GO:00016810; P:transport; IEA.
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O7MGU7;

O1-MAR-2004 (TEMBLrel. 26,

O1-MAR-2004 (TEMBLrel. 26,

O1-MAR-2004 (TEMBLrel. 26,
                                                                                                                                                                                                                                                                            PRINTS; PRO0344; BCTRLSENSOR.

PRODOM; PD000039; Response reg; 1.

PROSITE; PS50109; HIS KIN; 1.

PROSITE; PS50283; NA_SOLUT_SYMP_3; 1.

PROSITE; PS50110; RESPONSE REGULATORY; 1.

Kinase; Phosphorylation; Sensory transduction; Transferase.

SEQUENCE 1156 AA; 128283 MW; 4DB3AB41A0B1A4FB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR003594; ATPbind_ATPase.
InterPro; IPR004358; Bact sens pr_C.
InterPro; IPR005467; His Kinase.
InterPro; IPR003661; His kina N.
InterPro; IPR001734; Na/Solut_Symport.
InterPro; IPR001799; Response_reg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF02518; HATPase c; 1.
Pfam; PF00512; HisKA; 1.
Pfam; PF00072; Response reg; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chen C.-Y., Wu K.-M., Chang Y.-C., Chang C.-H., Tsai H.-C., Liao T.-L., Liu Y.-M., Chen H.-J., Shen A.B.-T., Li J.-C., Shao C.-P., Lee C.-T., Hor L.-T., Tsai S.-F.; "Comparative genome analysis of Vibrio vulnificus, a marine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vibrio vulnificus (strain Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genome Res. 13:2577-2587(2003).
-!- SIMILARITY: Contains 1 histidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=196600;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=VV3129;
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; Sensor histidine
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57 65
PSLSERLQSASFVGTPLPENENISLYQSRVTVGELEMLASRFVGRNRVKNAPAHYWSQQR
568 576
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                                                                                                                                                                                    October 13, 2004 13:25 (zara371.key)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                kinase.
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Last sequence update)
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578

ETLLPNQQAPSTLIRHTERVLAGVFGASSAKLVLTSALQGRNMQLEEVA

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1 match found in sequence: q86p24; RE22501p (Fragment). (from "claim36uni.pep") TOIG of: q86p24 check: 8357
                                                                                                                                                                                                                                                                                                                                                                                                     408
                                                                                                                                                                                                                                                                                                                                                                                                                                                   348
                                                                                                                                                                                                                                                                           l match found in sequence:
  q8dcz5; Signal transduct
  (from "claim36uni.pep")
  TOIG of: q8dcz5 check:
RRP RN COC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Found using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q86P24 Length: 972 October 13, 2
ound using 'claim36' (zara371.key)
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Q86P24;
01-JUN-2003
         STRAIN=CMCP6;
Rhee J.H., Ki
Choy H.E.;
                                                                                                                                            QBDCZ5 PRELIMINARY; PRT; 1143 AA.
QBDCZ5;
QBDCZ5;
Q1-MAR-2003 (TrEMBLrel. 23, Created)
Q1-MAR-2003 (TrEMBLrel. 23, Last sequence update)
Q1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Signal transduction histidine kinase.
OrderedLocusNames=VV11242;
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Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J.,

Stapleton M., Chavez C., Dorsett V., Dresnek D., Farfan D., Frise E.,

Champe M., Chavez C., Dorsett V., Dresnek D., Farfan D., Frise E.,

George R., Gonzalez M., Guarin H., Krommiller B., Li P., Liao G.,

Miranda A., Mungall C.J., Nunoo J., Pacleb J., Paragas V., Park S.,

Patel S., Phouanenavong S., Wan K., Yu C., Lewis S.E., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases. EMBL; BT00352; AA039526.1; -. FlyBase; FBgn0038721; CG6718. InterPro; IPR007632; DUF590.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RE22501p (Fragment).
ORFNames=CG16718;
                                                                                                       Bacteria; Proteobacteria;
Vibrionaceae; Vibrio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQÜENCE 972 AA; 112363 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
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01-MAR-2004 (TrEMBLrel.
                                                          SEQUENCE FROM N.A.
                                                                                     NCBI_TaxID=672;
                                                                                                                                     Vibrio vulnificus.
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                                                                                                                                                                                                                                                                                                          transduction histidine kinase.
                           Kim S.Y.,
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                                                                                                                                                                                                                                                                             2146 from: 1
                           Chung
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24,
                                                                                                                    Gammaproteobacteria; Vibrionales;
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                           s.s.,
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Last annotation update)
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                             Kin
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                           J.J.,
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                             Moon
                             Y.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Check:
                           Jeong
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565
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R GO; GO:0016020; C:membrane; IEA.

R GO; GO:0005524; F:ATP binding; IEA.

R GO; GO:0003677; F:DNA binding; IEA.

R GO; GO:0003677; F:NA binding; IEA.

R GO; GO:0016301; F:kinase activity; IEA.

R GO; GO:0016740; F:transferase activity; IEA.

R GO; GO:0015215; F:transporter activity; IEA.

R GO; GO:000155; F:two-component response regulator activity; GO; GO:0000155; F:two-component sensor molecule activity; GO; GO:0000155; F:two-component sensor; IEA.

R GO; GO:0000105; F:two-component signal transduction system of the component sequence of the component sequence of the component sensor of the com
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Pfam; PF02518; HATPase_C; 1.
Pfam; PF00512; HisKA; \(\bar{1}\).
Pfam; PF00072; Response_reg; 1.
Pfam; PF000744; BCTRLSENSOR.
PRINTS; PR00344; BCTRLSENSOR.
PRODOm; PD000039; Response_reg; 1.
SMART; SM00387; HATPase_C; 1.
SMART; SM00388; HisKA; 1.
SMART; SM00091; PAS; 1.
SMART; SM000448; REC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                 QBIN71
QBIN71;
                                                                                                                                       ORFNames=CG16718;
Drosophila melanogaster (Fruit fly).
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transferase.
SEQUENCE 1
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Submitted (DEC-2002) to the EM
-!- SIMILARITY: Contains 1 his
EMBL; AE016801; AA009698.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                            CG16718-PB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS50109; HIS KIN; 1.
PROSITE; PS50123; NA SOLUT SYMP 3; 1.
PROSITE; PS50110; RESPONSE REGULATORY; 1.
Complete proteome; Kinase; Phosphorylation; Sensory transduction;
                                                                              NCBI_TaxID=7227;
SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   [nterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PSLSERLQSASFVGTPLPENENISLYQSRVTVGELEMLASRFVGRNRVKNAFAHYWSQQR
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RA Golder C., Gabbielian A.E., Garg N.S., Gelbart W.M., Glasser K., RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Aharris M.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J., RA Harris M.L., Havey D., Heiman T.J., Wei M.H., Ibegwam C., RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., Alali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Liang Y., Lin X., Mattei B., McIntosh T.C., McIeod M.P., McPherson D., Liu X., Mattei B., McIntosh T.C., McLod M.P., McPherson D., R., McIntosh T.C., Mollod M.P., McPherson D., RA, Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A., RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M., Ra Melson D.R., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G., Ra Reinert K., Remington K., Saunders R.D., Scheeler F., Smith T., Ra Shue B.C., Siden-Kiamos I., Sinpson M., Skupski M.P., Smith T., Ra Spier E., Spradling A.C., Stapleton M., Strong R., Sun E., Svirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X., Wang Z.Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Weinsen S.M., WoodageT, Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yah R.F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L., Thou X., Molds G.M., Weinsenbach J., The genome sequence of Drosophila melanogaster."; Shi X., Science 287:2185-2195(2000).
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A Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
A George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
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A Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
B sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
B Barandon R.C., Rogers Y.H., Blazej R.G., Champe M., Pfeiffer B.D.,
A Barid J.F., Agbayani A., An H.J., Andrews-Pfannkoch C., Baldwin D.,
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A Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
A Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
A Cherry J.M., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
Dourbin K.J., Evangelista C.C., Ferraz C., Ferriara S., Fleischmann W.,
A Borley C., Schmann W.,
Charley J.M., Carlon A. G., Galhart W.,
Classor Y.
                                                                                                                                                                     MEDLINE=2246069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Misra S., Crosby M.A., Mungall C.J., Millburn G.H., Prochnik S.E..
Smith C.D., Tupy J.L., Whitfied E.J., Bayraktaroglu L., Berman B.P
Bettencourt B.R., Celniker S.E., de Grey A.D., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Celniker S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Halpern A., Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A., George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson C.R., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J., Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Venter C. Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M., "Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence.", Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
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Kaminker J.S., Bergman C.M., Kronmiller
Patel S., Frise E., Wheeler D.A., Lewis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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Ashburner M., Celniker S.
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                                                                                     "Annotation of the Drosophila melanogaster euchromatic genome: a
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3:RESEARCH0083-RESEARCH0083 (2002)
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1 match found in sequence:
q8lb06; Putative glucan
(from "claim36uni.pep")
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Jound using
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Q8LB06;
01-OCT-2002
                                                                         GO; GO:0004553; F:hydrolase activity, hydrolyzing O-glycosyl GO; GO:0005975; P:carbohydrate metabolism; IEA.
InterPro; IPR000490; Glyco.hydro 17.
Pfam; PF0032; Glyco.hydro 1.
PROSITE; PS00587; GLYCOSYL HYDROL F17; UNKNOWN 1.
SEQUENCE 460 AA; 50643 MW; DF3361006601162C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                      Genome
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MEDLINE=22088475; PubMed=12093376;
Haas B.J., Volfovsky N., Town C.D., Tro
Feldmann K.A., Flavell R.B., White O.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; edicotyledons; core e
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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01-MAR-2004 (TrEMBLrel. 26, Last annotation
Putative glucan endo-1-3-beta-glucosidase.
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EMBL, AE003727; AAN13804.1;

FlyBase; FBgn0038721; CG1671

InterPro; IPR007633; DUF590.

Pfam; PF04547; DUF590; 1.
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SEQUENCE FROM N.A.
FLYBASE;
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O., Salzberg
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                                                                                                                                                                                                                                                                                                                                                        Lu Y.-P.,
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IWCVVAKGANWTQLGDALSYACSQGNNTCDPIQRGGPCQKPDLTVLHASYAFSSYWAQFR

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386
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     HACTER SOOO SOON
                                                                                                                                                                                                                                                                                                                                                      t match found in sequence:
    q91k41;    Beta-1,3-glucanse.
    (from "claim36uni.pep")
    TOIG of: q91k41 check: 217
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q8lby4 ; Beta-1,3-glucanase,
(from "claim36uni.pep")
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Cound using 'claim36'
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                                 Beta-1,3-glucanase.
Arabidopsis thaliana (Mouse-ear cress).
Arabidopsis thaliana (Mouse-ear cress).
Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
surosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                      Q9LK41;
Q9LK41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF
PROSITE;
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Q8LBY4;
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01-OCT-2000 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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MEDLINE=22088475; PubMed=12093376;

Haas B.J., Volfovsky N., Town C.D., Troukhan M., Ale:
Feldmann K.A., Flavell R.B., White O., Salzberg S.L.,
Feldliength messenger RNA sequences greatly improve
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Arabidopsis thaliana (Mouse-ear cress).
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosida
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01-OCT-2002 (TrEMBLrel. 22,
01-MAR-2004 (TrEMBLrel. 26,
NCBI_TaxID=3702; [1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (MAR-2002) to the EMBL/GenBank/DDBJ EMBL; AY086926; AAM64490.1; -. HSSP; P15737; 1CHS.
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PROSITE; PR00323; GLYCOSYL_HYDROL_F17; UNKNOWN 1.

SEQUENCE 476 AA; 52087 MW; 22DDEEA17FA96EB3 CRC64;
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Feldmann K.;
                                                                                                                                                                                                                                                                                                                                       ; Bera-i, "claim36uni pep")
of o9lk41 check: 2170
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436 444
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26,
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Found using
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Q9SFW1; 17EMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.

Lin X., Kaul S., Town C.D., Benito M.-I., Crea

Ronning C.M., Koo H., Fujii C.Y., Utterback T.

Bowman C.L., White O., Nierman W.C., Fraser C.

Submitted (JAN-2001) to the EMBL/GenBank/DDBJ

EMBL; AC012395, AAF20214.1; -.

HSSP; P12257; 1AQ0
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudiotyledons; core e
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Kaneko T., Kato T., Sato S.,
Submitted (JUL-1999) to the:
EMBL; APO00377; BAB01853.1;
HSSP; P15737; 1GHS.
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Sequence features of
TAC and BAC clones.";
                                                                                                                                                                                                               GO; GO:0004553; F:hydrolase activity, hydrolyzing GO; GO:0005975; P:carbohydrate metabolism; IEA. InterPro; IPR000490; Glyco_hydro_17. Pfam; PF00332; Glyco_hydro_17; 1. PROSITE; PS00587; GLYCOSYL_HYDROL_F17; UNKNOWN 1. SEQUENCE 440 AA; 48538 MW; FC8B6A3B3B4B8D93 CF
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MEDLINE=20363099; PubMed=10907853;
Nakamura Y.;
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; Putative beta-1,3-glucanase
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  KTEYKESLPAPENNDLYKGKIWCVGNNTCDPIQRGGPCQKPDLTVLHASYAFSSYWAQFR
                                                                                                                                   Length: 440
ing 'claim36'
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Res. 7:217-221(2000).
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                                                                                                                                   October 13, (zara371.ke
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EMBL/GenBank/DDBJ databases.
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R.M.;
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Barnstead
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KIGGTCSFNGLATQTIKDPSYGRCEFPSVTL

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430
                                                                                                                                                                                                                                                                                 370
                l match found in sequence:
    q9vdv4 ; CG16718-PA (LD10322p).
    (from "claim36uni pep")
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(from "claim36uni.pep")
TOIG of: q9srt4 check: 8766
                                                                                                                                                                                                                                                                                                                                                                                                                                   Found
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Q9SRT4;
01-MAY-2000
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yamada K., Chan M.M., Chang C.H., Dale J.M., Hsuan V.W., Lee J.M. Onodera C.S., Quach H.L., Tang C., Toriumi M., Wong C., Wu H.C., Yu G., Yuan S., Carninci P., Chen H., Cheuk R., Hayashizaki Y., Ishida J., Jones T., Kamidya A., Kawai J., Kim C.J., Narusaka M., Nguyen M., Palm C.J., Sakurai T., Satou M., Seki M., Shinn P., Southwick A., Tripp M.G., Wu T., Shinozaki K., Davis R.W., Ecker Theologis A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Putative glucan endo-1-3-beta-glucosidase (Putative glucanase) (Putative glycosyl hydrolasae), Putative glycosyl hydrolasae), Name=R1307320/T1B9 1; Arabidopsis thaliana (Mouse-ear cress).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.

Lin X., Kaul S., Town C.D., Benito M.-I., Creasy T.H., Haas Lin X., Kaul S., Town C.D., Utterback T.R., Barnstead Ronning C.M., Koo H., Fujii C.Y., Utterback T.R., Barnstead Bowman C.L., White O., Nierman W.C., Fraser C.M.; Submitted (JAN-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core evenosids II; Brassicales; Brassicaceae; Arabidopsis.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AC009853; AAF02143.1; -.
EMBL; AK118068; BAC42699.1; -.
EMBL; BT005678; AAO64098.1; -.
EMBL; BT005678; AAO64098.1; -.
EMBL; BT005678; F:hydrolase activity; IEA.
GO; GO:0016787; F:hydrolase activity, hydrolyzing
GO; GO:0004553; F:hydrolase activity, hydrolyzing
GO; GO:0005975; P:carbohydrate metabolism; IEA.
InterPro; IPR000490; Glyco_hydro_17.
Pfam; PF00332; Glyco_hydro_17; 1.
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Seki M., Iida K., Satou M., Sakurai T., Akiyama K., Ishida
Nakajima M., Bnju A., Kamiya A., Narusaka M., Carninci P.,
Hayashizaki Y., Shinozaki K.;
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PROSITE; PS00587; GLY
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420
428
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                                                                                                                                                                                                                                                                                                                                                                                                                                   October 13, 2004 13:25 (zara371.key)
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1097 from:
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Kawai
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RX MEDLINE 2019606; PubMed=10731132;
RX Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
RA Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
Sutton G.G., Wortman J.R., Yandell M.D., Zlang Q., Chen L.X.,
RA Sutton R.C., Rogers Y.H., Blazej R.G., Champe M., Ffeiffer B.D.,
RA Barlon R.C., Rogers Y.H., Blazej R.G., Champe M., Ffeiffer B.D.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Borkova D., Botchan M.R., Bouck J., Bavengari D., Bolshakov S.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Bavengari L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davengort L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davengort L.B., Davies P.,
RA Cherry J.M., Berman D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heinan T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heinan T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Kaaft C., Kravitz S., Kulp D., Lai Z.,
RA Jalaii M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Lisko P., Lei Y., Levitsky A.A., Li J., Li J., Li J., Li J., Li Z., Liang Y., Lin X.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
Mount S.M., Moy M., Murphy B., Murphy L., Muny D., Lai Z.,
RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Nelson D.L.,
RA Nelson D.R., Nelson K.A., Sunders R.D., Scheeler F., Shen H.,
RA Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Sheng X.H., Modage T., Wor
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Q9VDV4;
Q9VDV4;
01-MAY-2000
01-MAY-2000
05-JUL-2004
                                                         SEQUENCE FROM N.A.
MEDLINE=22426070; PubMed=12537573;
Kaminker J.S., Bergman C.M., Kronm
Wheeler D.A.,
                                                                                                                                           Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A. George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson C.R. Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E. Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Ver Weinstock G., Scherer S.E., Myers E.W., Glbbs R.A., Rubin G.M. "Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence.";

Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygol
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
               The transposable
                                                                                                                                                                                                                                                                                         MEDLINE=22426065; PubMed=12537568; Celniker S.E., Wheeler D.A., Kronm
                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                   Patel S.,
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                 niker S.E. elements
                                                   Wheeler D.A.,
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Last
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Kronmiller B., Carlson J.W., Hal
L., Dugan S.P., Frise B., Hodgson
Muzny D.M., Nelson C.J
                                                                      Kronmiller
                   the
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                 Drosophila
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                                                                      Carlson
                   melanogaster
                                                     Rubin
                                                     G.M., S
                                                                      Svirskas
                                                                                                                                                                                                                                                                                           Halpern
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Number of sequences searched:
Number of sequence hits:
Number of separate matches:
Number of sequence hits saved:
                                                                                                                                                                                                                                          511
                                                                                                            Times:
                                                                                                                                                                                                                                                                                                451
                                                                                                                                                                                                                                                                                                                                                                                                                                             Found using
                                                                                                                                                                                                                                                                                                                                                                                     Q9VDV4 Length: 1075 October 13, 2004 13:25 ound using 'claim36' (zara371.key)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.

STRAIN-Berkeley;

Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J.,

Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J.,

Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J.,

Stapleton M., Frise E.,

George R., Gonzalez M., Guarin H., Kronmiller B., Li P., Liao G.,

Miranda A., Mungall C.J., Nunco J., Pacleb J., Paragas V., Park S.,

Patel S., Phouanenavong S., Wan K., Yu C., Lewis S.E., Rubin G.M.,

Celniker S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AE003727; AAF55685.1; -.
EMBL; BT010299; AAQ23517.1; -.
IntAct; Q9VDV4; -.
F1yBase; Esgn0038721; CG16718.
InterPro; IPR007632; DUF590.
Pfam; PF04547; DUF590.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIINE=22426069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Misra S., Crosby M.A., Mungall C.J., Millburn G.H., Prochnik S.E.,
Smith C.D., Tupy J.L., Whitfied B.J., Bayraktaroglu L., Berman B.P.,
Bettencourt B.R., Celniker S.E., de Grey A.D., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Geriome Biol. 3:RESEARCH0084-RESEARCH0084(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (MAR-2004) to the EMBL/GenBank/DDBJ databases.
[7]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A. FLYBASE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Annotation of the Drosophila melanogaster euchromatic genome: a systematic review.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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[5]
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501 509
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                                                                                                                                                 Search Statistics
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p72824 ; Hypothetical protein
(from "claim36pt.pep")
  Run mode
Time to start comparison
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Lected sequence key from "zara371.
claim36 (AA) ID claim36 AA preli
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P72824;
01-FEB-1997
01-FEB-1997
                                        MEDLINE-97061201; PubMed-8905231; Kaneko T., Sato S., Kotani H., Tanaka A., Asamizu E., Nakamura Kaneko T., Sato S., Kotani H., Tanaka A., Asamizu E., Kimura T., Miyajina N., Hirosawa M., Sugiura M., Sasamoto S., Kimura T., Hosouchi T., Mateuno A., Muraki A., Nakazaki N., Naruo K., Okum Shimpo S., Takeuchi C., Wada T., Watanabe A., Yamada M., Yasuda Tabata S.;
"Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PC66803. II. Sequence determination of the entire genome and assignment of potential protein-coding regions."; DNA Res. 3:109-136(1996).
                                                                                                                                                                   01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-CCT-2003 (TrEMBLrel. 25, Last annotation updat
Hypothetical protein sll1200.
                                                                                                                                    Synechocystis sp. (strain PCC 6803).
Bacteria; Cyanobacteria; Chroococcales;
                                                                                                       SEQUENCE FROM N.A.
                                                                                                                           NCBI_TaxID=1148;
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M., Yasuda
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                                                                                                                                                                                 q86p24 ; RE22501p (Fragment).
(from "claim36spt.pep")
TOIG of: q86p24 check: 8357
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(from "claim36spt.pep")
TOIG of: q06913 check: 219 from: 1
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Q86P24;
Q86P24;
01-JUN-2003
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01-OCT-2003
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Q06913;
Q1-NOV-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beta-1,3-glucanase homologue (Fragment).

Brassica napus (Rape).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids;

Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids;

Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids;
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EMBL; X69889; CAA49515.1; -.
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346 354
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m; PF00332; Glyco_hydro_17; 1.
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    q8dcz5; Signal transduction histidine kinase.
    (from "claim36spt.pep")
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
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                                                                                                                                                                                                                                                                                                                       EMBL;
                                                                                                                                                                                                                                                                                                                                                                                               Rhee J.H., Kim S.Y., Chung S.S., Kim J.J., Moon Y.H., Jeong Choy H.E.;
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CG16718.
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Vibrionaceae; Vibrio.
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                Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases EMBL; AE016801; AA009698.1; -... GO; GO:0016020; C:membrane; IEA. GO; GO:0005524; FiATP binding; IEA. GO; GO:0003677; F:DNA binding; IEA. GO; GO:0016301; F:kinase activity; IEA. GO; GO:0015215; F:two-component response regulator acti GO; GO:0005215; F:two-component sensor molecule activit GO; GO:000155; F:two-component sensor molecule activit GO; GO:0000157; F:two-component sensor molecule activit GO; GO:0000159; F:two-component signal transduction sys InterPro; IPR00359; ATPbind ATPase.

InterPro; IPR00359; ATPbind ATPase.
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                                               InterPro;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LELWKRYSAEI THRWDLTGFDVHEEHPRPQYLARLEHI PPTRVDYVTNI
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BT003522; AAO39526.1; -.
Pro; IPR007632; DUF590.
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                                                                                                                             regulator activity; IEA plecule activity; IEA.
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                                                                             IEA.
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1 match found in sequence: q8in71; CG16718-PB.
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(from "claim36ept.pep")
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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams M.D., Celniker S.E., Li P.W., Hookins R.A., Galle R.F.,
RA Amanatides P.G., Scherer S.E., Li P.W., Hookins R.A., Galle R.F.,
RA George R.A., Lewis S.E., Randell M.D., Zhang Q., Chen L.X.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Wan K.H., Doyle C., Baxeer E.G., Helt G., Nelson C.R., Gabor G.L.,
RA Wan K.H., Doyle C., Baxeer E.G., Helt G., Nelson C.R., Gabor G.L.,
RA Wan K.H., Doyle C., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Ghoriy J.M., Cawley S., Dahlke C., Davenport L.B., Center A., Chandra I.,
RA Ge Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Posler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
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Pfam; PF00512; HAIPAse c; 1.

Pfam; PF00512; response_reg; 1.

Pfam; PF00072; response_reg; 1.

PRINTS; PR00344; BCTRLSENSOR.

ProDom; PD000039; Response_reg; 1.

SMART; SM00387; HAIPAse_c; 1.

SMART; SM00388; HiskA; I.

SMART; SM00448; REC; 1.

SMART; SM00448; REC; 1.

SMART; SM0048; REC; 1.

PROSITE; PS50109; HIS KIN; 1.

PROSITE; PS50109; HIS KIN; 1.

PROSITE; PS50110; RESPONSE_REGULATORY; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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CG16718-PB.
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01-MAR-2003
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SEQUENCE 1143 AA; 126643 MW; 85F47E27E3B2D621 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20196006; PubMed=10731132;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM
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555 563
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IPR001734; Na/solut_symport.
IPR001789; Response_reg.
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(TrEMBLrel.
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Last annotation update)
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362
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  match
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RA Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
RA Reinert K., Remington K., Saunders R.D., Scheeler F., Shen H.,
RA Shue B.C., Siden-Klamos I., Simpson M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Syirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Wang Z.Y., Waasarman D.A., Weinstock G.M., Weissenbach J.,
RA Williams S.M., WoodsgeT, Worley K.C., Wu D., Yang S., Yao Q.A., Ye J.,
RA Yeh R.F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
RA Zheng X.H., Zhong F.N., Zhong W., Zhou X., Smith H.O.,
RA Gibbs R.A., Myers E.M., Rubin G.M., Venter J.C.;
RT "The genome sequence of Drosophila melanogaster.";
RI Science 287:2185-2195(2000).
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Banzon J., An H., Baldwin D., Banzon J., Beeson K.Y., Busam D.A.,
RA Carlson J.W., Center A., Champe M., Davenport L.B., Dietz S.M.,
Dodson K., Dorsett V., Doup L.E., Doyle C., Dresnek D., Farfan D.,
Regriera S., Frise E., Galle R.F., Garg N.S., George R.A.,
Gonzalez M., Houck J., Hoskins R.A., Hostin D., Howland T.J.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
Ra Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
RA Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
Ra Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
Ra Ibegwam C., Jalali M., Kruse D., Li P., Mattei B., Moshrefi A.,
Ra Ibegwam C., Jalali M., Kruse D., Jalali M., Hoskins R., Tochin C., Roshrefi A.,
Ra Ibegwam C., Jalali M., Kruse D., Jalali M., Houck M., Houck M., Houck M., Hoskins R., Tochin C., Roshrefi A.,
Ra Ibegwam C., Jalali M., Houck M., Ho
                                                                                                                                                                                                                                                                                                                                                                                                      Q8IN71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misra S., Crosby M.A., Matthews B.B., Bayraktaroglu L., Campbell K Hradecky P., Huang Y., Kaminker J.S., Prochnik S.E., Smith C.D., Tupy J.L., Bergman G., Berman B., Carlson J.W., Celniker S.E., Clamp M., Drysdale R., Emmert D., Frise E., de Grey A., Harris N., Kronmiller B., Marshall B., Millburn G., Richter J., Russo S., Searle S.M.J., Smith E., Shu S., Smutniak F., Whitfield E., Ashburner M., Gelbart W.M., Rubin G.M., Mungall C.J., Lewis S.E.; "Annotation of Drosophila melanogaster genome."; Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AE003727; AAN13804.1; -. FlyBase; FBgn0038721; CG16718. InterPro; IPR007632; DUF590.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
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Celniker S.E., Ada
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Adams M.D., Celniker
Submitted (MAR-2000)
  found
                                                                                                                                                                                                    VPVKDICQSGNTNITMCPLCDWCNFWDLKETCNYAKVTYLIDNPSTVFFAVFMSFWATLF
352 36
                                                                                                                                            LELWKRYSAEITHRWDLTGFDVHEEHPRPQYLARLEHIPPTRVDYVTNI
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ing 'claim36'
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ij
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sequence:
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Amanatides P.G., Brandon
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EMBL/GenBank/DDBJ databases.
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370
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  (from "claim36spt.pep")
TOIG of: q8lb06 check: 8289 from: 1 to: 460
  RREAD OCCUPANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        match found in sequence: q81by4; Beta-1,3-glucana
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(from "claim36spt.pep")
                                                                                                                                                                                                                                                                                                                                                                                        OBLBY4;
OBLBY4;
01-OCT-2002
01-OCT-2002
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Q8LB06;
01-OCT-2002
01-OCT-2002
                                                                                                                                                                                                                                                                   EMBL; AY087496; AAM65039.1; -
GO; GO:0004553; F:hydrolase activity, hydrolyzing O-gl:
GO; GO:005975; P:carbohydrate metabolism; IEA.
InterPro; IPR000490; Glyco_hydro_17.
Pfam; PF00332; Glyco_hydro_17; 1.
PROSITE; PS00587; GLYCOSYL_HYDROL_F17; 1.
SEQUENCE 460 AA; 50643 MW; DF3361006601162C CRC64;
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;

eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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01-0CT-2002 (TrEMBLrel. 22, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
Putative glucan endo-1-3-beta-glucosidase.
                                                                         Haas B.J., Volfovsky N. Feldmann K.A., Flavell "Full-length messenger annotation.";
                                                                                                                                                                                                                                                           Spermatophyta; Magnoliophyta; eurosids II; Brassicales; Bras
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SEQUENCE FROM N.A.
                                                   Genome Biol. 0:0-0(2002).
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                                                                                                                                                                                    SEQUENCE FROM N.A.
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, Volfovsky N., Tow.
K.A., Flavell R.B.,
K.A., Flaver RNA F
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White O.,
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D., Salzberg S.
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  (from "claim36spt.pep")
TOIG of: q9lk41 check: 217
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Q9LK41;
01-OCT-2000
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                                                                                                                                                                                                                                                                                    EMBL; AP000377; BAB01853.1; -.
HSSP; P15737; 1GHS.
GO: GO: 0004553; F: chydrolase activity, hydrolyzing O-glycosyl.
GO; GO: 0004553; F: carbohydrate metabolism; IEA.
InterPro; IFR000490; Glyco hydro_17.
Pfam; PF00332; Glyco hydro_17.
PROSITE; PS00587; GLYCOSYL HYDROL F17; 1.
SEQUENCE 476 AA; 52170 MW; 9B36D1BA6109B46E CRC64;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core e
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Kaneko T., Kato T.,
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Pfam; PF00332; Glyco_hydro_17; 1.

PROSITE; PS00587; GLYCOSYL HYDROL F17; 1.

SEQUENCE 476 AA; 52087 MW; 22DDEEA17FA96EB3 CRC64;
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NA Res. 7:217-221(2000).
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(from "claim36spt.pep")
                            q9srt4 ; Putative glucan endo-1-3-beta-glucosidase
(from "claim36spt.pep")
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C Q9SRT4;
C Q9SRT4;
T 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
T 01-MAY-2000 (TrEMBLrel. 25, Last annotation update)
T 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
P PUTATIVE Glucan endo-1-3-beta-glucosidase (Putative beta precursor) (Putative glycosyl hydrolase).
E precursor) (Putative glycosyl hydrolase).
N F2103.3 OR AT3G07320/T1B9_1 OR AT3G07320.
S Arabidopsis thaliana (Mouse-ear cress).
C Spermatophyta; Magnoliophyta; Embryophyta; Trac Spermatophyta; Magnoliophyta; eudicotyledons; core eudic cursids II; Brassicales; Brassicaceae; Arabidopsis.
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EMBL; AC012395; AAR20214.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GO; GO:0004553; F:hydrolase activity, hydrolyzing GO; GO:0005975; P:carbohydrate metabolism; IEA. InterPro; IER000490; Glyco_hydro_17. Pfam; PF00332; Glyco_hydro_17; 1. PROSITE; PS00587; GLYCOSYL HYDROL F17; 1. SEQUENCE 440 AA; 48538 MW; FC8B6A3B3B4B8D93 CR
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; edicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
STRAIN=CV. Columbia;
STRAIN=CV. Columbia;
Lin X., Kaul S., Town C.D., Benito M., Creasy T.H., Haas B.,
Ronning C.M., Koo H., Fujii C.Y., Utterback T.R., Barnstead M.E.,
Bowman C.L., White O., Nierman W.C., Fraser C.M.;
Bowman C.L., White O., Nierman W.C., Fraser C.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAY-2000
01-JUN-2003
                                                                                                                                                                                                                                    NCBI_TaxID=3702;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T1B9.1
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                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             found in sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KTEYKESLPAPENNDLYKGKIWCVGNNTCDPIQRGGPCQKPDLTVLHASYAFSSYWAQFR
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ing 'claim36'
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) (TrEMBLrel. 13,
) (TrEMBLrel. 24,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          check: 8766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       October 13, 2 (zara371.key)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       from: 1
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Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Check:
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                                                                                                                                                                                                                                                                                                       eudicots; rosids;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         beta-1,3-glucanase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        400
                                                                                                                                                                                                                                                                                                                                Tracheophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence.";
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430
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                q9vdv4 ; CG16718 protein.
(from "claim36spt.pep")
TOIG of: q9vdv4 check: 1097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Found using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             match found in sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q9SRT4 Length: 460
Cound using 'claim36'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Arabidopsis Open Reading Frame (ORF) Clones.";
Submitted (MAR-2003) to the EMBL/GenBank/DDBJ database
EMBL; AC009853; AAF02143.1; -.
EMBL; AK118068; BAC42699.1; -.
EMBL; BT005678; AA064098.1; -.
EMBL; BT005678; AA064098.1; -.
HSSP; P12257; 1AQO.
GO; GO:0004553; F:hydrolase activity, hydrolyzing O-c
GO; GO:0016787; F:hydrolase activity; IEA.
GO; GO:0015975; P:caarbohydrate metabolism; IEA.
InterPro; IPR000490; Glyco_hydro_17.
Pfam; PF00332; Glyco_hydro_17; 1.
PROSITE; PS00587; GLYCOSYL_HYDROL_F17; 1.
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Yamada K., Chan M.M., Chang C.H., D

Yamada K., Chan H.L., Tang C.,

Onodera C.S., Quach H.L., Tang C.,

Yu G., Yuan S., Carninci P., Chen H

Ishida J., Jones T., Kamiya A., Kaw

Nguyen M., Palm C.J., Sakurai T., S

Southwick A., Tripp M.G., Wu T., Sh
STRAIN=Berkeley;
STRAIN=Berkeley;
STRAIN=Berkeley;
STRAIN=Berkeley;
MEDLINE=20196006; PubMed=10731132;
Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
Adams M.D., Celniker S.E., Li P.W., Hoskins R.A., Galle R.P.,
Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.P.,
Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Henderson S.N.,
George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
Sutton G.G., Wortman J.R., Yandell M.D., Champe M., Pfeiffer B.D.,
Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
Brandon R.C., Rogers Y.-H.C., Blazej R.G., Nelson C.R., Miklos G.L.G.,
Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
                                                                                                                                                                                                                                                                                                                          Q9VDV4 PRELI
Q9VDV4; O1-MAY-2000 (TYEW
01-MAY-2000 (TYEW
01-JUN-2003 (TYEW
CG1671B protein.
CG1671B.
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SEQUENCE
                                                                                                                                                                                                                                             Drosophila melanogaster (Fruit fly).
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Bekaryota; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=cv. Columbia;
Seki M., Iida K., Satou
Nakajima M., Enju A., Ka
Hayashizaki Y., Shinozak
                                                                                                                                                                                                                         Ephydroidea; Dro
NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Arabidopsis thaliana full-length cDNA.";
Submitted (NOV-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                        SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IWCVVAKGANWTQLGDALSYACSQGNNTCDPIQRGGPCQKPDLTVLHASYAPSSYWAQFR
420
428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KIGGTCSFNGLATQTIKDPSYGRCEFPSVTL
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                                                                                                                                                                                                                                                                                                                                                                 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                     (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       50603 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chang C.H., L., Tang C., Toriumi m., ... Tang C., Toriumi m., ... Tang C., Toriumi m., Hayashizan. Inci P., Chen H., Cheuk R., Hayashizan. Ramiya A., Kawai J., Kim C.J., Narusaka M., Kamiya A., Kawai J., Kahim C.J., Satou M., Seki M., Shinn P., Satou M., Seki M., Shinn P., Satou M., Satou M., Seki M., Shinn P., Satou M., Seki M., Bcker
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Kamiya A., Narusaka
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13,
24,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     from: 1
                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                       Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       D2061EAA09C0F85F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to: 1075
                                                                                                                                                                                                                                                                                                                                                                                                                                              1075
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ka M., Carninci P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                              ð
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Kawai J.,
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Britch Project S., Sprach S., Stapleton M., Strong G., Zhao Q., Zheng L., RMBL, Aboug T., Diffs 20. Drosophila melanogaster.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Found
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q9VDV4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     V4 Length: 1075 using 'claim36'
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Pfam; PF04547; DUF590;
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501 509
                                                                                                                                                                                                                                                                              LELWKRYSAEITHRWDLTGFDVHEEHPRPQYLARLEHIPPTRVDYVTNI
f sequences searched:
f sequence hits:
f separate matches:
f sequence hits saved:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1075
                                                                                                          00:00:00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA;
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                                                                                                                                                                        Search Statistics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     123934 MW;
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    0111
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                                                                                                                                                         submitted to the EMBL Data Library, December 1992
A;Description: The anther specific protein encoded by the Brassica napus and Arabidopsis thaliana A6 gene exhibits homology to beta-1,3-glucanases.
A;Reference number: S31612
A;Accession: S31612
A;Molecule type: mRNA
A;Residues: 1-139 <HIR>
A;Residues: 1-139 <HIR>
                                                                                                                                                                                                                                                                                                                                                  F1;S31612 - beta-1,3-glucanase homolog (clone A20) - rape (fragment) C;Species: Brassica napus (rape) C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 17-Nov-2000 C;Accession: S31612 R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, D.; Hodge, R.; Paul, W.; Smartt, S.; Draper, J.; Scott, R;Hird, D.; Worrall, R;Hird, R
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831612; TOIG of: 831612 check: 219 from: 1
(from "claim36pir.pep")
TOIG of: 831612 check: 219 from: 1 to: 139
                                                                     A;Cross-references: EMBL:X69889; NID:g17733; PID:g17734
A;Experimental source: clone A20
C;Superfamily: beta-1,3-glucanase
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A; Variety: PCC 6803

A; Variety: PCC 6803

C; Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 08-Oct-1999

C; Accession: S74688

R; Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.;
Miyajima, N.; Hirosawa, M.; Sugiura, M.; Sasamoto, S.; Kimura, T.; Hosouchi, T.; Matsuno, A.; Muraki, A.; Nakazaki, N.; Naruo, K.; Okumura, S.; Shimpo, S.
Takeuchi, C.; Wada, T.; Watanabe, A.; Yamada, M.; Yasuda, M.; Tabata, S.
DNA Res. 3, 109-136, 1996

DNA Res. 3, 109-136, 1996

A; Fittle: Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. PCC6803. II. Sequence determination of the entire genome assignment of potential protein-coding regions.

A; Reference number: S74322; MUID:97061201; PMID:8905231

A, Accession: S74688
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s74688; TOIG of: s74688

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A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-391 <KAN>
A;Cross-references: EMBL:D90901; GB:AB001339; NID:g1651897; PIDN:BAA16839.1;
PID:d1017572; PID:g1651913
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aab17079
                                                                                                                                                                                            aab17079 ; Mdm/hdm antagonist peptide sequence SEQ ID NO:135.
(from "claim36ags.pep")
TOIG of: aab17079 check: 5978 from: 1 to: 12
                                                                                                                                                                                                                    match found in sequence
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        04-MAY-2000
                                                                         AAB17079 standard; peptide; 12 AA
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                         WO200024782-A2
                                                          thrombosis;
                                                                                                                          Mdm/hdm antagonist peptide sequence SEQ ID NO:135
                                                                 cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor; vascular endothelial growth factor; matrix metalloproteinase;
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. match found in sequence:
aab17080 ; Mdm/hdm antagonist peptide sequence SEQ
(from "claim36ags.pep")
TOIG of: aab17080 check: 6151 from: 1 to: 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC Fc domain, partmacologically active peptides, and linkers. Where (II) is:

CC (X1)a-F1-(X2)b, where: F1 = an Fc domain; X1 and X2 = are each

CC independently selected from -(L1)c-F1-(L1)d-F2-(L1)d-F2, -(L1)c-F1-

CC (L2)d-P2-(L3)e-P3, or -(L1)c-F2-(L2)d-P2-(L3)e-P3-(L4)f-P4 where P1, P2,

CC P3, and P4 = are each independently sequences of pharmacologically active

CC peptides; L1, L2, L3, and L4 = are each independently linkers; and a, b,

CC c, d, e, and f = are each independently 0 or 1, provided that at least 1

CC of a and b is 1. The composition can have cytostatic, antiasthmatic,

CC thrombolytic and immunosuppressive activities. DNAs, vectors and host

CC cells from the present invention can be used for producing pharmaceutical

CC compositions. The compositions are useful for treating cancer, asthma,

CC thrombosis, or autoimmune diseases. The use of an Fc domain (rather than

CC a Fab domain) can provide a longer half-life or incorporate functions

CC anability placental transfer. AAA69443 to AAA69526 and AAB16955 to

CC AAB18003 represent nucleotide and amino acid sequences used in the

CC exemplification of the present invention
                                                                                                                                                                               autoimmune disease; cytostatic; antiasthmatic; thrombolytic; VESF; immunosuppressive; EPO; TPO; CTLA4; mimetic; IL-1; TNF; antagonist inhibitor; erythropoietin; thrombopoietin; interleukin 1; cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 12 AA;
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22-OCT-1999;
                                                                                                          Synthetic
                                                                                                                                                                                                                                                     Modified peptide; therapeutic agent; fusion; Fc domain; cancer;
                                                                                                                                                                                                                                                                                             Mdm/hdm antagonist peptide sequence SEQ ID NO:136.
                                                                                                                                                                                                                                                                                                                                                                       AAB17080;
                                                                                                                                                                                                                                                                                                                                                                                                         AAB17080 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes composition of matter (I) comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Feige
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-OCT-1999;
                                                                                                                                              thrombosis; pharmaceutical.
                                                                                                                                                              vascular endothelial growth factor; matrix metalloproteinase; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MPRFMDYWEGLN
3 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     composition of matter comprising an Fc domain and pharmacologically peptides, useful for treating cancer and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                39; Page 241; 608pp; English.
                                                                                                                                                                                                                                                                                                                                (first entry)
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99US-00428082.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US025044
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                                                                                                                                                                                                                                                                                                                                                                                                       peptide;
                                                                                                                                                                                                                                                                                                                                                                                                           12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NO:136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Check: 5978
                                                                                                                                                                                                                      antagonist;
                                                                                                                                                                                                                        MP;
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25-OCT-1999; 04-MAY-2000

99WO-US025044.

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aab17081 ; Mdm/hdm antagonist peptide sequence SEQ ID NO:137.
  (from "claim36ags.pep")
TOIG of: aab17081 check: 5993 from: 1 to: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Found using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes composition of matter (I) comprising an CC (XI)a-F1-(XZ)b, where: F1 = an Fc domain; XI and X2 = are each CC (IX)a-F1-(XZ)b, where: F1 = an Fc domain; XI and X2 = are each CC (IX)a-F2-(IX)b-P3, or -(IX)c-P1, -(IX)c-P1-(IZ)d-P2, -(IX)c-P1-CC (IZ)d-P2-(IX)e-P3, or -(IX)c-P1-(IZ)d-P2-(IX)e-P3 (IX)f-P4 where P1, P2, CR (IX)a-P2-(IX)e-P3, and IX = are each independently linkers; and a, b, CC c, d, e, and f = are each independently 0 or 1, provided that at least 1 CC of a and b is 1. The composition can have cytostatic, antiasthmatic, CC thrombolytic and immunosuppressive activities. DNAs, vectors and host CC cells from the present invention can be used for producing pharmaceutical CC compositions. The compositions are useful for treating cancer, asthma, CC thrombosis, or autoimmune diseases. The use of an Fc domain (rather than CC such as Fc receptor binding, protein A binding, complement fixation, and Dossibly placental transfer. AAA69443 to AAA69526 and AAB16955 to AAB18003 represent nucleotide and amino acid sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  match found in sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB17080 Length: 12
ound using 'claim36'
                                                       04-MAY-2000
                                                                                                                              Synthetic
                                                                                                                                                                                                      Modified peptide; therapeutic agent; fusion; Fc domain; cancer; autoimmune disease; cytostatic; antiasthmatic; thrombolytic; VEGF; immunosuppressive; EPO; TPO; CTLA4; mimetic; II-1; TNF; antagonist; MMP; inhibitor; erythropoietin; thrombopoietin; interleukin 1; cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor;
                                                                                                                                                                                                                                                                                                                                                                31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel composition of matter comprising an Fc domain and pharmacologically active peptides, useful for treating cancer and autoimmune diseases.
               25-OCT-1999;
                                                                                                                                                                      thrombosis;
                                                                                                                                                                                                                                                                                                                      Mdm/hdm antagonist peptide sequence SEQ ID NO:137.
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB17081 standard; peptide; 12 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 39; Page 242; 608pp; English
                                                                                           WO200024782-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-OCT-1998;
22-OCT-1999;
                                                                                                                                                                                      cyrotoxic T cell lymphocyte antigen 4; tumour necrosis factor;
vascular endothelial growth factor; matrix metalloproteinase; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AMGE-) AMGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VQNFIDYWTQQF
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                                                                                                                                                                      pharmaceutical.
                                                                                                                                                                                                                                                                                                                                                              (first entry)
                 99WO-US025044.
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99US-00428082.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   October 13, (zara371.key)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cheetham J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2004 13:39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boone TC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Type: P
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PC domain, pharmacologically active peptides, and linkers. Where (I) is:

(X) A.Fl-(X2)b, where: Fl = an Fc domain; Xl and X2 = are each independently selected from -(L1)c-Pl, -(L1)c-Pl-(L2)d-P2, -(L2)d-P2, -(L1)c-Pl-(L2)d-P2, -(L2)d-P2, -(L2)d-P2, -(L1)c-Pl-(L2)d-P2, -(L2)d-P2, -(
Sequence 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-350702/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-OCT-1998;
22-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 active peptides, useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Feige U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  composition of matter comprising an Fc domain and pharmacologically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          39; Page 242; 608pp; English.
   Ā
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention describes composition of matter (I) comprising pharmacologically active peptides, and linkers. Where (I)
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99US-00428082
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer and autoimmune diseases.
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AABI7081 Length: 12 Found using 'claim36' October 13, 2004 13:39 (zara371.key) Type; P Check: 5993

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TGPAFTHYWATF
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l match found in sequence:
aabi7082; Mdm/hdm antagonist peptide sequence SEQ ID
(from "claim36egs:pep")
TOIG of: aabi7082 check: 9093 from: 1 to: 15 AAB17082 standard; peptide; 15

31-OCT-2000 (first entry)

Mdm/hdm antagonist peptide sequence SEQ ID NO:138.

cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor; vascular endothelial growth factor; matrix metalloproteinase; autoimmune disease; cytostatic; antiasthmatic; immunosuppressive; EPO; TPO; CTLA4; mimetic; iinhibitor; erythropoietin; thrombopoietin; into 25-OCT-1999; Synthetic 04-MAY-2000 WO200024782-A2 thrombosis; Modified peptide; therapeutic agent; fusion; Fc domain; pharmaceutical. 99WO-US025044. interleukin IL-1; TNF; thrombolytic; VEGF; antagonist;

23-OCT-1998;

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match found in sequence:
aab17083; Mdm/hdm antagonist peptide sequence SEQ
(from "claim36esp.pep")
TOIG of: aab17083 check: 9428 from: 1 to: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes composition of matter (I) comprising an CC Fc domain, pharmacologically active peptides, and linkers. Where (I) is: CC (X1)a-F1-(X2)b, where: F1 = an Fc domain; X1 and X2 = are each ccc independently selected from -(L1)c-P1, -(L1)c-P1-(L2)d-P2, -(L1)c-P1-(L2)d-P2, -(L1)c-P1-(L2)d-P2, -(L1)c-P1-(L2)d-P2, -(L1)c-P1-(L2)d-P2, -(L1)c-P1-(L2)d-P2, -(L1)f-P4 where P1, P2, CC (L2)d-P2-(L3)e-P3, or -(L1)c-P1-(L2)d-P2-(L3)e-P3-(L4)f-P4 where P1, P2, CC (P3, and P4 = are each independently sequences of pharmacologically active CC c, d, e, and f = are each independently 0 or 1, provided that at least 1 cf a and b is 1. The composition can have cycostatic, antiasthmatic, CC compositions and invention can be used for producing pharmaceutical CC compositions. The compositions are useful for treating cancer, asthma, CC compositions or autoimune diseases. The use of an Fc domain (rather than a Fab domain) can provide a longer half-life or incorporate functions CC such as Fc receptor binding, protein A binding, complement fixation, and CC composition of the present invention acid sequences used in the CC chalbago 3 represent nucleotide and amino acid sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB17082 Length: 15
Cound using 'claim36'
                                                                                                                                                                                                                  inhibitor; erythropoietin; thrombopoietin; interleukin 1;
cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor;
vascular endothelial growth factor; matrix metalloproteinase; asthma;
                                                                                                                                                                                                                                                                    Modified peptide; therapeutic agent; fusion; Fc domain; cancer; autoimmune disease; cytostatic; antiasthmatic; thrombolytic; VEGF; ammunosuppressive; EPO; TPO; CTLA4; mimetic; IL-1; TNF; antagonist;
                                                                                                                                                                                                                                                                                                                                                                                          31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB17083 standard; peptide; 15
23-OCT-1998;
22-OCT-1999;
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                                                    25-OCT-1999;
                                                                                                                               WO200024782-A2
                                                                                                                                                                  Synthetic
                                                                                                                                                                                                  thrombosis; pharmaceutical.
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6 14
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                                                                                                                                                                                                                                                                                                                                                   antagonist peptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 242; 608pp; English
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                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
98US-0105371P.
99US-00428082.
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                                                      99WO-US025044
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                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO:139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Check: 9093
                                                                                                                                                                                                                                                                            MMP;
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aab17084 ; Mdm/hdm antagonist peptide sequence SEQ
  (from "claim36ags.pep")
TOIG of: aab17084 check: 8833 from: 1 to: 15
AAB17083 Length: 15 Found using 'claim36'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            match found in sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cells from the present invention can be used for producing pharmaceutical compositions. The compositions are useful for treating cancer, asthma, thrombosis, or autoimmune diseases. The use of an Fc domain (rather than a Fab domain) can provide a longer half-life or incorporate functions such as Fc receptor binding, protein A binding, complement fixation, and possibly placental transfer. AAA6943 to AAA69526 and AAB16955 to AAB18003 represent nucleotide and amino acid sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P3, and P4 = are each independently sequences of pharmacologically active peptides; L1, L2, L3, and L4 = are each independently linkers; and a, b, c, d, e, and f = are each independently 0 or 1, provided that at least 1 of a and b is 1. The composition can have cytostatic, antiasthmatic, thrombolytic and immunosuppressive activities. DNAs, vectors and host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes composition of matter (I) comprising a FC domain, pharmacologically active peptides, and linkers. Where (I) (X1)a-F1-(X2)b, where: F1 = an FC domain; X1 and X2 = are each independently selected from -(L1)c-P1, -(L1)c-P1-(L2)d-P2, -(L1)c-P1-(L2)d-P2, -(L3)e-P3-(L4)f-P4 where P1, (L2)d-P2-(L3)e-P3-(L4)f-P4 where P1, (L2)d-P3-(L3)e-P3-(L4)f-P4 where P1, (L2)d-P3-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4)e-P4-(L4
                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              autoimmune disease; cytostatic; antiasthmatic; thrombolytic; VEGF; immunosuppressive; EPO; TPO; CTLA4; mimetic; IL-1; TNF; antagonist inhibitor; erythropoietin; thrombopoietin; interleukin 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB17084 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                      thrombosis; pharmaceutical.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytotoxic T cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modified peptide; therapeutic agent; fusion;
                                                                                                                                                                                                                                                         WO200024782-A2
                                                                                                                                                                                                                                                                                                                                                                                                                     cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor;
vascular endothelial growth factor; matrix metalloproteinase; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mdm/hdm antagonist peptide sequence
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6 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 243; 608pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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23-OCT-1998; 22-OCT-1999;

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l match found in sequence:
aag07078; krabidopsis thaliana protein fragment SEQ
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Found using 'claim36'
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                                                                                                             New sequence determined DNA fragments (SDFs) from different plant species, e.g. corn, rice or Arabidopsis thaliana, useful as promoters, protein coding sequences, untranslated regions, or as 3' termination
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Claim 19;

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The present sequence is a

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The present sequence is a putative protein fragment from Arabidopsis thaliama. Its coding sequence was isolated by carrying out RT-PCR on all of the mRNA obtained from the plant, and creating a cDNA library which could then be sequenced, allowing the putative protein sequence(s) to be obtained. This sequence may be useful for protein identification and for aiding in the elucidation of signal transduction and metabolic pathways. Its coding sequence has a use in the control of gene expression as a promoter, coding sequence, 3'UTR or termination sequence, for controlling the behaviour of a gene within the chromosome, as a tool for use in genetic mapping, including a use in hybridisation assays, for recognition or isolation of similar DNA fragments, or for the identification of a particular organism
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Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.
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ID AAG38987 standard; protein; 493 AA.

XX
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DT 18-OCT-2000 (first entry)

XX
DT Arabidopsis thaliana protein fragment SEQ ID NO: 48177.

XX
Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter;

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W Explosis thaliana.

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PN Ep1033405-A2.

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      The present sequence is a putative protein fragment from Arabidopsis thaliana. Its coding sequence was isolated by carrying out RT-PCR on all of the mRNA obtained from the plant, and creating a cDNA library which could then be sequenced, allowing the putative protein sequence(s) to be obtained. This sequence may be useful for protein identification and for aiding in the elucidation of signal transduction and metabolic pathways. Its coding sequence has a use in the control of gene expression as a promoter, coding sequence, 3'UTR or termination sequence, for controlling the behaviour of a gene within the chromosome, as a tool for use in genetic mapping, including a use in hybridisation assays, for recognition or isolation of similar DNA fragments, or for the identification of a
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430
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26-JAN-2000;
27-JAN-2000;
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27-JAN-2000;
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                                                                                                                                                                                                  Sequence 460 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New sequence determined DNA fragments (SDFs) from different plant species, e.g. corn, rice or Arabidopsis thaliana, useful as promoters, protein coding sequences, untranslated regions, or as 3' termination
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N-PSDB; AAC45900.
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Zheng L, Dum
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. match found in sequence:
aag38989;; Arabidopsis thaliana protein fragment SEQ
(from "claim36ags.pep")
TOIG of: aag38989 check: 1002 from: 1 to: 384
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06-MAY-1999
06-MAY-1999
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21-MAY-1999
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05-MAR-1999

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20-APR-1999

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99US-0127462P

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99US-0130510P

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99US-0132484P

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354
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                                                                                                                                                                                                                      l match found in sequence:
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(from "claim16ags pep")
TOIG of: aag52473 check: 8156
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N-PSDB; AAC45900.
                  Arabidopsis thaliana protein fragment SEQ
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aag52474; Arabidopsis thaliana
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     25-FEB-1999;
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                                                                                                                                                                                               thaliana. Its coding sequence was isolated by carrying out RT-PCR on all of the mRNA obtained from the plant, and creating a cDNA library which could then be sequenced, allowing the putative protein sequence(s) to be obtained. This sequence may be useful for protein identification and for aiding in the elucidation of signal transduction and metabolic pathways. Its coding sequence has a use in the control of gene expression as a promoter, coding sequence, 3'UTR or termination sequence, for controlling the behaviour of a gene within the chromosome, as a tool for use in genetic mapping, including a use in hybridisation assays, for recognition or isolation of similar DNA fragments, or for the identification of a particular organism
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NQSSQCYFNGLARETTTNPGNERCKFPSVTL

match found in sequence:
aag52475; Arabidopsis thaliana
(from "claim36ags pep")
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18-OCT-2000 (first entry)

Arabidopsis thaliana protein fragment SEQ ID ö 66708

Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence

Arabidopsis thaliana

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Zheng L, I
The present sequence is a putative protein fragment from Arabidopsis thaliana. Its coding sequence was isolated by carrying out RT-PCR on all of the mRNA obtained from the plant, and creating a cDNA library which could then be sequenced, allowing the putative protein sequence(s) to be obtained. This sequence may be useful for protein identification and for aiding in the elucidation of signal transduction and metabolic pathways. Its coding sequence has a use in the control of gene expression as a promoter, coding sequence, 3'UTR or termination sequence, for controlling the behaviour of a gene within the chromosome, as a tool for use in
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N-PSDB;
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      aaw37170 ; Human oncogenic protein MDM2 binding peptide 1.
(from "claim36ags.pep")
TOIG of: aaw37170 check: 5978 from: 1 to: 12
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This peptide is capable of binding to an oncogenic protein MDM2 (especially human DM2). The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2,p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can
                                                                                                                                                                                                                                                          Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
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                                                                                                                                                                                                                            Claim 5; Page 41; 45pp; English.
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07-APR-1997;
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binding; vi
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Chene P,
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AAW37170 Length: 12 Found using 'claim36'
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     CC (especially human DM2). The MDM2 binding peptides can specifically CC (inhibit or block the binding of MDM2 to the human p53 protein, in vitro CC inhubit or block the binding of MDM2 to the human p53 protein, in vitro CC or in vivo. Inhibiting the interaction between the p53 and MDM2 can CC induce growth arrest or apoptosis in tumour cells comprising a wild-type CC molecules that bind to MDM2 and to identify and design inhibitors of CC especially MDM2, diagnose disease by measuring levels of MDM2 in blood of CC cancer and leukaemia patients and for treatment or prevention of disease CC involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by CC injection. By interfering with MDM2/p53 interaction, the peptides can CC which mimic the MDM2 binding site in p53, have a significantly greater CC blocking activity compared with wild-type p53
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aaw37172 ; Human oncogenic protein MDM2 binding peptide
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TOIG of: aaw37172 che
                                                                                                                                                                                                                                                                             Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly great blocking activity compared with wild-type p53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interaction; cancer;
                                                                                                                                                                                                                                                                                                interaction with p53 and viral infections
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A,

REMDYWEGL

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aaw37179 ; Human oncogenic protein MDM2 binding peptide derivative
  (from "claim36ags.pep")
TOIG of: aaw37179 check: 3427 from: 1 to: 9
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                      This peptide is capable of binding to an oncogenic protein MDM2 (cespecially human DM2). The MDM2 binding peptides can specifically CC inhibit or block the binding of MDM2 to the human p53 protein, in vitro cor in vivo. Inhibiting the interaction between the p53 and MDM2 can cC induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify complecules that bind to MDM2 and to identify and design inhibitors of CC especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease cinvolving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by compared with MDM2/p53 interaction, the peptides can cativate p53 function and accumulation in normal cells. The peptides can chicking activity compared with wild-type p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         match found in sequence:
Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                          Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW37179 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                                                                                                                Disclosure; Page 8; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                      Lane D, Boettger V, Garcia-Echeverria C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-JUL-1996;
07-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human oncogenic protein MDM2 binding peptide derivative 7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MDM2; oncogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NOVS ) NOVARTIS AG.
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6 14
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ng 'claim36'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis; binding;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96GB-00014197.
97GB-00007041.
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(zara371.key)
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Chene P, Furet P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   p53; human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Pickeley S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inhibition;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
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AAW37179 Length: 9 Found using 'claim36'

October 13, 2004 13:39 (zara371.key)

Type: P

Check:

3427

Sequence 12 AA;

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aaw37182 ; Human oncogenic protein MDM2 binding N-acylated peptide derivative
  (from "claim36ags.pep")
TOIG of: aaw37182 check: 5993 from: 1 to: 12
                         This is a N-acylated peptide derivative capable of binding to a human concogenic protein MDM2. The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of m0lecules that bind to MDM2 and to identify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
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                                                                                                                                                                                                                                                                                                                                                               Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lane D, Boettger V,
Garcia-Echeverria C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW37182 standard; peptide; 12 AA
                                                                                                                                                                                                                                                                                                                               Example 1; Page 18; 45pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-100996/09.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumour;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human oncogenic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          iomo sapiens.
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97GB-00007041.
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ng; viral infection.
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1 match found in sequence:
    aaw37183 ; Human oncogenic protein MDM2 binding N-acylated peptide derivative
    (from "claim35ags.pep")
    TOIG of: aaw37183 check: 5978 from: 1 to: 12
Found using
        This is a N-acylated peptide derivative capable of binding to a human CC oncogenic protein MDM2. The MDM2 binding peptides can specifically CC inhibit or block the binding of MDM2 to the human p53 protein, in vitro CC or in vivo. Inhibiting the interaction between the p53 and MDM2 can CC induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify CC molecules that bind to MDM2 and to identify and design inhibitors of CC (MDM2/p53 binding. They may also be used to purify binding partners CC especially MDM2, diagnose disease by measuring levels of MDM2 in blood of CC cancer and leukaemia patients and for treatment or prevention of disease CC involving p53/MDM2 interactions, especially tumours and viral infections. CC injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides CC which mimic the MDM2 binding site in p53, have a significantly greater CC blocking activity compared with wild-type p53
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Cound using 'claim36'
                                                                                                                                                                                                                                                                                                                      Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Lane D, Boettger V, Garcia-Echeverria C,
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(CANC-) CANCER RES (
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4 12
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                                                                                                                                                                                                                                                                                           Page 19;
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97GB-00007041.
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is; binding; viral infection.
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12
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                                                                                                                                                                                                                                                                                       45pp;
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Chene P, Furet P;
                                                                                                                                                                                                                                                                                           English.
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l match found in sequence:
aaw37191; Human oncogenic protein MDM2 binding N-acylated
(from "claim36egs.pep")
TOIG of: aaw37191 check: 6151 from: 1 to: 12
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This is a N-acylated peptide derivative capable of binding to a human CC concegenic protein MDM2. The MDM2 binding peptides can specifically controlled to the human p53 protein, in vitro CC inhibit or block the binding of MDM2 to the human p53 protein, in vitro CC or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and mon-elevated levels of MDM2. The peptides may be used to identify CC molecules that bind to MDM2 and to identify and design inhibitors of CC especially MDM2, diagnose disease by measuring levels of MDM2 in blood of CC especially MDM2 interactions, especially tumours and viral infections. The peptides can be administered masally, rectally, orally or by CC injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides CC which mimic the MDM2 binding site in p53, have a significantly greater
                                                                                                                                                                                                                                                                                    Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                  Lane D, Boettger V, Garcia-Echeverria C,
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07-APR-1997;
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                                                                                                                                                                                                                                                        Example 1; Page 19; 45pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human oncogenic
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3 11
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CANCER RES CAMPAIGN TECHNOLOGY.
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97GB-00007041
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Chene P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "N-terminal acetyl"
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AAW37191 Length: 12 Found using 'claim36'
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aaw37192 ; Human oncogenic protein MDM2 binding N-acylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (from "claim36ags.pep":
TOIG of: aaw37192 chec
                                         This is a N-acylated peptide derivative capable of binding to a human oncogenic protein MDM2. The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease
                                                                                                                                                                                                                                        Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                    Lane D, Boettger V, Garcia-Echeverria C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW37192 standard; peptide; 15 AA
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involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can
                                                                                                                                                                                                                                                                                                         WPI; 1998-100996/09.
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                                                                                                                                                                                                               Example 1; Page 19;
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07-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MDM2; oncogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW37192;
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                                                                                                                                                                                                                                                                                                                                                                               (NOVS ) NOVARTIS AG. (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ncogenic protein; p53; human; inhibit
diagnosis; binding; viral infection.
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97GB-00007041.
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                                                                                                                                                                                                             45pp; English.
                                                                                                                                                                                                                                                                                                                                      Chene P, Furet P;
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                                                                                                                                                                                                                                                                                                                                                    Picksley S,
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1 match found in sequence:
        aaw37193 ; Human oncogenic protein MDM2 binding N-acylated peptide derivative
  (from "claim36ags.pep")
  TOIG of: aaw37193 check: 9428 from: 1 to: 15
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This is a N-acylated peptide derivative capable of binding to a human oncogenic protein MDM2. The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections.
                                                                                                                                                                                                                      Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
                                                                                                                                                                                                                                                                                                                     Lane D, Boettger V,
Garcia-Echeverria C,
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07-APR-1997;
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                                                                                                                                                                                          Example 1; Page 19; 45pp; English.
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ng 'claim36'
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97GB-00007041.
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is; binding; viral infection.
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Chene P, Furet P;
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aaw37194 ; Human oncogenic protein MDM2 binding N-acylated peptide
(from "claim36ags.pep")
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
This is a N-acylated peptide derivative capable of binding to a human oncogenic protein MDM2. The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of
                                                                                                                                                                Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
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                                                                                                                                                                                                                                                     Garcia-Echeverria
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g 'claim36'
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CANCER RES CAMPAIGN TECHNOLOGY.
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diagnosis; binding; v:
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cheverria C,
                                                                                                                                          Page 19;
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97GB-00007041.
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                                                                                                                                         45pp; English.
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Chene P, Furet P;
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viral infection
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                                                                                                                                                                                                                                                                  Hochkeppel H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
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match found in sequence:
aaw37195 ; Human oncogeni
  (from "claim36ags.pep")
  TOIG of: aaw37195 check
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                                                                                                                                                                                       Lane D, Boettger
Garcia-Echeverria
                                                                                                                                                                                                                                                                                 04-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                                                                                                                                    WPI; 1998-100996/09.
                                                                                                                                                                                                                     (NOVS )
                                                                                                                                                                                                                                                                                                                                                                                                                           MDM2; oncogenic
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2 10
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                                                                                                                                                                                                                                                                                                                                                                                                               ncogenic protein;
diagnosis; bindir
                                                                                                                                                                                                                     NOVARTIS AG.
CANCER RES CAMPAIGN TECHNOLOGY.
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97GB-00007041.
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                                                                                                                                                                                                                                                                                                                                             /note=
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binding; v:
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Chene P,
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.key)
                                                                                                                                                                                                                                                                                                                                                                                                                  viral infection.
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Furet
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                                                                                                                                                                                                                                                                                                                                             amide"
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This is a C-amidated peptide derivative capable of binding to a human oncogenic protein MDM2. The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of

Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections

Example 1; Page

20; 45pp; English. binding agents.

ful in, e.g. identifying

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match found in sequence:
aaw37196 ; Human oncogenic
(from "claim36ags.pep")
TOIG of: aaw37196 check:
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This is a C-amidated peptide derivative capable of binding to a human oncogenic protein MDM2. The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of
                                                                                                                                                                                Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
                                                                                                                                                                                                                                             WPI; 1998-100996/09.
                                                                                                                                                                                                                                                                           Lane D, Boettger V, Garcia-Echeverria C,
                                                                                                                                                                                                                                                                                                                                                                   05-JUL-1996;
07-APR-1997;
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                                                                                                                                                     Example 1; Page 20; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                04-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
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97GB-00007041.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide;
                                                                                                                                                                                                                                                                                                                       CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             binding; viral infection.
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                                                                                                                                                                                                                                                                           Boettger Chene P,
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Furet
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ret P;
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match found in sequence:
aaw37204; Human oncogenic protein MDM2 binding biotinylated peptide derivativ
(from "claim36ags.pep")
TOIG of: aaw37204 check: 1571 from: 1 to: 28
AAW37196 Length: 12 Found using 'claim36'
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                                                                                                             Compounds binding to MDM2 protein and inhibit useful in, e.g. diagnosis and treatment of car and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
                                                                                                                                                                                    Lane D, Boettger
Garcia-Echeverria
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07-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis;
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heverria C,
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97GB-00007041.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein MDM2 binding biotinylated
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                                                                                                                                                                                                                                                                                                                                                                'note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      binding;
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Chene P, Furet P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      viral infection; biotinylation
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                                                                                                                                                                                               Hochkeppe1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interaction;
                                                                                                                            interaction with p53 and viral infections
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         derivative 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer;
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This is a biotinylated peptide derivative capable of binding to a human oncogenic protein MDM2. The MDM2 binding peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of

3; Page

21; 45pp; English.

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aaw37205; Human oncogenic protein MDM2 binding biotinylated peptide derivativ
(from "claim36age.pep")
TOIG of: aaw37205 check: 8233 from: 1 to: 31
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                                                     useful in, e.g. diagnosis and t
and identifying binding agents.
                                                                                        Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections
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07-APR-1997;
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CANCER RES C
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97GB-00007041
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                                                                                                                                                                                                                                                                                                                                   CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                       Boettger .
Chene P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "beta-Alanine"
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Example 3; Page 21; 45pp; English

Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.

WPI; 1998-100996/09.

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AAW37205 Length: 31 Found using 'claim36'
match found in sequence:
aaw37216 ; Human oncogenic protein MDM2 binding peptide derivative
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TOIG of: aaw37216 chec
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                                                                                  Lane D, Boettger V, Garcia-Echeverria C,
                                                                                                                                                                            05-JUL-1996;
07-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 MDM2; oncogenic protein; p53; human; inhibition; tumour; diagnosis; binding; viral infection.
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                                                                                                                                                                                                                                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                    Boettger A, Picks
Chene P, Furet P;
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                                                                                                    Hochkeppel
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match found in sequence:
aaw37220 ; MDM2 binding peptide unique phage insert sequence
(from "claim36ags.pep")
TOIG of: aaw37220 check: 5978 from: 1 to: 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
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                                                                                                                                Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
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07-APR-1997;
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                                                                              Example 8; Page 30; 45pp; English
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                                                                                                                                                                                                                                                                                                         Boettger
Chene P,
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Furet
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inhibition;
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                                                                                                                                                                      and
                                                                                                                                                                                             interaction with p53
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This is a unique phage insert identified by phage display. 7

sequence of the The MDM2 binding

MDM2 binding peptides and

peptide

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match found in sequence:
aaw37221; MDM2 binding p
  (from "claim36ags.pep")
TOIG of: aaw37221 check
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cc derivatives are capable of binding to the human oncogenic protein MDM2. CT These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to cells comparison to the peptides of MDM2, p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring converted to the peptides of tancer and leukaemia patients and for creatment or prevention of disease involving p53/MDM2 interactions, cespecially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 contraction, the peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53 contractions.
  This is a unique phage insert sequence of the MDM2 binding peptide identified by phage display. The MDM2 binding peptides and their derivatives are capable of binding to the human oncogenic protein MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumou
                                                                                                                                                                             Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.
                                                                                                                                                                                                                                                                                                    Lane D, Boettger V,
Garcia-Echeverria C,
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pb3 protein,
e p53 and wr
                                                                                                                                         Page 30; 45pp;
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97GB-00007041.
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l match found in sequence:
aaw37222; MDM2 binding peptide unique phage insert sequence
(from "claim36age.pep")
TOIG of: aaw37222 check: 5993 from: 1 to: 12
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07-APR-1997;
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Chene P, Furet P;
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                                                                                                                                                                                                                                                                                       Picksley S,
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This is a unique phage insert sequence of the MDM2 binding peptide identified by phage display. The MDM2 binding peptides and their derivatives are capable of binding to the human oncogenic protein MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring

by measuring also be used 8; Page

30;

45pp; English.

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match found in sequence:
aaw37223; MDM2 binding peptide unique phage insert sequence
(from "claim36egs.pep")
TOIG of: aaw37223 check: 9093 from: 1 to: 15
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This is a unique phage insert sequence of the MDM2 binding peptide identified by phage display. The MDM2 binding peptides and their derivatives are capable of binding to the human oncogenic procesin MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to gentify and design inhibitors of MDM2,553 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53
                                                                                                                                                                                                                                                                                                            Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
                                                                                                                                                                                                                                                                                                                                                                                                                        Lane D, Boettger V, Garcia-Echeverria C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-1996;
07-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tumour;
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                                                                                                                                                                                                                                                                             Example 8; Page
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using 'claim36'
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diagnosis; bindi
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97GB-00007041.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inhibition;
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l match found in sequence:
aaw37224; MDM2 binding peptide unique phage insert
(from "claim36ags.pep")
TOIG of: aaw37224 check: 9428 from: 1 to: 15
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         This is a unique phage insert sequence of the MDM2 binding peptide CC identified by phage display. The MDM2 binding peptides and their CC identified by phage display. The MDM2 binding peptides and their CC derivatives are capable of binding to the human oncogenic protein MDM2. These peptides can specifically inhibit or block the binding of MDM2 to CC the human p53 protein, in vitro or in vivo. Inhibiting the interaction CC between the p53 protein, in vitro or in vivo. Inhibiting the interaction CC cells comprising a wild-type p53 and non-elevated levels of MDM2. The CC geptides may be used to identify molecules that bind to MDM2 and to CC identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring CC treatment or prevention of disease involving p53/MDM2 interactions, CC especially tumours and viral infections. The peptides can be administered nesally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
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                                                                                                                                                                                                                                                                                                                                                                   Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, hav a significantly greater blocking activity compared with wild-type p53
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07-APR-1997;
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CANCER RE
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97GB-00007041.
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Chene P,
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l match found in sequence:
aaw37225; MDM2 binding peptide unique phage insert sequence
(from "claim36ags.pep")
TOIG of: aaw37225 check: 8833 from: 1 to: 15
                      CC This is a unique phage insert sequence of the MDM2 binding peptide CC identified by phage display. The MDM2 binding peptides and their CC derivatives are capable of binding to the human oncogenic protein MDM2. CC These peptides can specifically inhibit or block the binding of MDM2 to CC the human p53 protein, in vitro or in vivo. Inhibiting the interaction CC between the p53 and MDM2 can induce growth arrest or apoptosis in tumour Cc cells comprising a wild-type p53 and non-elevated levels of MDM2. The CC identify and design inhibitors of MDM2, p53 binding. They may also be used to identify molecules that bind to MDM2 and to CC identify binding partners especially MDM2, diagnose disease by measuring CC treatment or prevention of disease involving p53/MDM2 interactions, CC especially tumours and viral infections. The peptides can be administered constally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in commal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with wild-type p53
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Sequence 15 AA;
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07-APR-1997;
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g 'claim36'
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97GB-00007041
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Chene P, Furet P;
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AAW37225 Length: 15 Found using 'claim36'

October 13, (zara371.key)

2004 13:39

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Check:

8833

PAFSRFWSDLSAGAH
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aaw82322 ; p53 homologue TIP 12/1 peptide
  (from "claim36ags.pep")
TOIG of: aaw82322 check: 4400 from: 1
AXU
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aaw8230; p53 homologue TIP 12/1 peptide.
(from "claim36ags.pep")
TOIG of: aaw82320 check: 4400 from: 1
                                                                                                               match found in sequence:
                                                                                                                                                                                                                             AAW82320 Length: 19
ound using 'claim36'
                                                                                                                                                                                                                                                                                                                                           This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the
                                                                                                                                                                                                                                                                                Sequence 19 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2.
                             AAW82322 standard; peptide; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
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                                                                                                                                                                                                                                                                                                                             treatment of cancer, viral conditions or other conditions associated with
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(zara371.key)
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match found in sequence:
abb65993; Drosophila melanogaster polypeptide SEQ ID
(from "claim36ags.pep")
TOIG of: abb65993 check: 1097 from: 1 to: 1075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW82322 Length: 19 October 13, Found using 'claim36' (zara371.key)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows level p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 AA;
                                           WO200171042-A2
                                                                       Drosophila melanogaster
                                                                                                       pharmaceutical.
                                                                                                                                                 Drosophila melanogaster polypeptide SEQ ID NO 24771.
                                                                                                                                                                                 26-MAR-2002
                                                                                                                                                                                                                                            ABB65993 standard; protein; 1075 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig 1; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New substance with a mdm2 binding domain and coupling partner - useful for stabilising in cells without an efficient mdm2-mediated degradation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-OCT-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                    Drosophila; developmental biology; cell signalling; insecticide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        non-functional p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-609975/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9847919-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  p53 homologue TIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-FEB-1999
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7 15
                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                             2004 13:39
                                                                                                                                                                                                                                                                                                                                                                                                                                             Type:
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                                                                                                                                                                                                                                                                                                        NO 24771.
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1 match found in sequence:
abb73174; Mdm/hdm antagonist peptide SEQ ID NO:135.
(from "claim36ags.pep")
TOIG of: abb73174 check: 5978 from: 1 to: 12
451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABB65993 Length: 1075
Found using 'claim36' (
                                                                                erythropoletin; TPO; tumour necrosis factor alpha inhibitor; TWP; TNF-alpha inhibitor; interleukin I antagonist; TL-1 antagonist; TPO mimetic peptide; EPO mimetic peptide; EMP; VEGF antagonist; MMP inhibitor; antiinflammatory; antitumour; immunosuppressive; cytostatic; antirheumatic; antiarthritic; antidiabetic; ophthalmologi. antianaemic; antorectic; antiinfertility; haemostatic; dermatological; neuroprotective; inflammatory disease; autoimmune disease; tumour grocancer; rheumatoid arthritis; diabetic retinopathy; infertility; obes sleep disorder; neurological degenerative disease; anaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from brosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher enkaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL6176-ABL30511), expressed DNA sequences (ABL01840-ABL16175) and the encoded proteins (ABB57737-ABB72072). The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                             ABB73174;
                                                                                                                                                                                                                                                                                                                                                                                             ABB73174 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-MAR-2000; 2000US-0191637P
11-JUL-2000; 2000US-00614150
                                                                                                                                                                                                                                                                                   Mdm/hdm antagonist peptide SEQ ID NO:135.
                                                                                                                                                                                                                                                                                                                        05-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1075 AA;
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              Homo sapiens
                                                                                                                                                                                                                                                 Modified peptide; mimetic; Fc domain; fusion;
                                                                  thrombocytopaenia; metastatic tumour; systemic
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                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detection reagent for detecting 1000 or more for elucidating cell signaling and cell-cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Type: P
                                                                                                                                                                                                                                                 immunoglobulin G;
                                                                  lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Check: 1097
                                                                                                                                                          ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention
alling and
                                                                                                                                                                                                                                                 IgG;
                                                                                                      r growth; obesity;
                                                                                                                                                                                                                                                   EPO;
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neuroprotective activities. (I) can be used as a therapeutic or prophylactic agent as well as for screening purposes. (I) is useful for diagnosing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of interest, in a biological sample. Additionally, (I) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders, infertility, and neurological degenerative diseases. (I), comprising EPO-mimetic compounds are useful for treating disorders characterised by low red blood cell levels such as anaemia. The TPO-mimetic comprising compounds are useful for treating conditions that involve an existing megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet
                                           tumour which result in thrombocytopaenia, aplastic anaemia, metastatic tumour which result in thrombocytopaenia, systemic lupus erythematosus, and Fanconi's syndrome. ABB72403 to ABB73426 and ABL35695 to ABB35777 represent amino acid and nucleic acid sequences used in the exemplification of the present incoming sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                     multimers. (I) can have antiinflammatory, antitumour, immunosu cytostatic, antirheumatic, antiarthritic, antidabetic, ophtha antianaemic, anorectic, antiinfertility, haemostatic, dermatol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel vehicle-peptide molecule or its multimers useful for treating inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
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Sequence 12 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     39; Page 53; 176pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Liu C, Cheetham JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention describes a vehicle-peptide molecule (I) or
                                         the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunosuppressive, c, ophthalmological
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ABB73174 Length: 12 Found using 'claim36' October 13, (zara371.key 2004 13:39 Type: ъ Check: 5978

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MPRFMDYWEGIN
3 11
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abb73175 ; Mdm/hdm antagonist peptide SEQ
 (from "claim36ags.pep")
TOIG of: abb73175 check: 6151 from: 1 match found in sequence: . 6 ID NO:136.

ABB73175 standard; peptide; 12 Ä

W W W W W W X E X F X A X X I ABB73175;

05-APR-2002 (first entry)

Mdm/hdm antagonist peptide SEQ ID NO:136

Modified peptide; mimetic; Fc domain; fusion; immunoglobulin G; erythropoletin; TPO; tumour necrosis factor alpha inhibitor; TNF-alpha inhibitor; interleukin I antagonist; IL-1 antagonist; TPO mimetic peptide; EPO mimetic peptide; EMP; VEGF antagonist; MMP inhibitor; antiinflammatory; antitumour; immunosuppressive; cytostatic; antirheumatic; antiarthritic; antidiabetic; ophthalm ophthalmological; EPO;

antianaemic; anorectic; antiinfertility; haemostatic; dermatological;

Mdm/hdm antagonist peptide

SEQ ID NO:137.

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SKKKE
                                                                                                                                                                                                                                                                                                                          ABB73175 Length: 12 Found using 'claim36'
                                                                                                                                                                                                                                                                                                                                                                                          match found in sequence:
abb73176; Mdm/hdm antagonist peptide SEQ
(from "claim36ags.pep")
TOIG of: abb73176 check: 5993 from: 1
                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a vehicle-peptide molecule (I) or its computations (I) can have antiinflammatory, antitumour, immunosuppressive, correstatic, antirheumatic, antiarthritic, antidiabetic, ophthalmological, antianaemic, anorectic, antiinfertility, haemostatic, dermatological and neuroprotective activities. (I) can be used as a therapeutic or composing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of interest, as a part of diagnostic kit to detect the presence of their proteins of interest in a biological sample. Additionally, (I) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, infertility, and neurological sample. Additionally, (I) is useful for treating district compounds are useful for treating disorders, and neurological degenerative diseases. (I), comprising conditions are useful for treating disorders characterised by low red blood cell levels such as anaemia. The TPO-mimetic compounds are useful for treating conditions that involve an existing megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet thrombocytopaenia, aplastic anaemia, metastatic tumour which result in thrombocytopaenia, splastic anaemia, metastatic and the present amino acid and nucleic acid sequences used in the expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet and nucleic acid sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel vehicle-peptide molecule or its multimers useful for treating inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility.
05-APR-2002
                                                                            ABB73176 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuroprotective; inflammatory disease; autoimmune disease; tumour growth; cancer; rheumatoid arthritis; diabetic retinopathy; infertility; obesity; sleep disorder; neurological degenerative disease; anaemia;
                                        ABB73176
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                          Ā
(first entry)
                                                                                                                                                                                                                                                                                                                            October 13, 2004 13:39 (zara371.key)
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                                                                                                                                                                ID NO:137
                                                                                                                       12
                                                                                                                                                                                                                                                                                                                                               Type: P
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1 match found in sequence:

TGPÄFTHYWATÉ 4

ABB73176 Length: 12 Found using 'claim36'

October 13, (zara371.key)

2004 13:39

Type:

ש

Check: 5993

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cc multimers. (1) can have antiinflammatory, antidiabetic, ophthalmological, cc cytostatic, antirheumatic, antiarthritic, antidiabetic, ophthalmological, cc antianaemic, anorectic, antiarthritic, haemostatic, dermatological and neuroprotective activities. (1) can be used as a therapeutic or prophylactic agent as well as for screening purposes. (1) is useful for cd diagnosing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of compounds are useful compounds are useful continuated proteins of interest in a biological sample. Additionally, (1) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, cc infertility, and neurological degenerative diseases, tumour growth, cancer, cc infertility, and neurological degenerative diseases. (1), comprising EPO-cmimetic compounds are useful for treating disporders characterised by low cred blood cell levels such as anaemia. The TPO-mimetic comprising compounds are useful for treating disporders characterised by low cred blood cell levels such as anaemia. The TPO-mimetic comprising compounds are useful for treating disporders characterised by low cred blood cell levels such as anaemia. The TPO-mimetic comprising compounds are useful for treating disporders characterised by low cred blood cell levels such as anaemia. The TPO-mimetic comprising compounds are useful for treating disporders characterised by low red blood cell in thrombocytopaenia, systemic anaemia, metastatic deficiency, such as thrombocytopaenia, systemic lugus crythematosus, in the present inventor anaemia, metastatic crypresent maino acid and nucleic acid sequences used in the crypresent amino acid and nucleic acid sequences used in the
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Synthetic.
Sequence 12 AA;
                                                                                                                                                      represent amino exemplification
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abb73177; Mdm/hdm antagonist peptide SEQ ID NO:138. (from "claim36ags.pep")
TOIG of: abb73177 check: 9093 from: 1 to: 15
                                                    prophylactic agent as well as for screening purposes. (I) is useful for diagnosing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of interest, as a part of diagnostic kit to detect the presence of their proteins of interest in a biological sample. Additionally, (I) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders, infertility, and neurological degenerative diseases. (I), comprising EPO-mimetic compounds are useful for treating disorders characterised by low red blood cell levels such as anaemia. The TPO-mimetic comprising compounds are useful for treating conditions that involve an existing megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency, such as thrombocytopaenia, spetemic lupus erythematossus, and Fanconi's syndrome. ABB72403 to ABB73426 and ABL35695 to ABL35777 represent amino acid and nucleic acid sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a vehicle-peptide molecule (I) or its multimers. (I) can have antiinflammatory, antitumour, immunosuppressive, cytostatic, antirheumatic, antiarthritic, antiidabetic, ophthalmological antianaemic, anorectic, antiinfertility, haemostatic, dermatological and neuroprotective activities. (I) can be used as a therapeutic or prophylactic agent as well as for screening purposes. (I) is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel vehicle-peptide molecule or its multimers useful for treating inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified peptide; mimetic; Fc domain; fusion; immunoglobulin G; IgG; EPO; erythropoietin; TPO; tumour necrosis factor alpha inhibitor; TMF; TNF-alpha inhibitor; interleukin 1 antagonist; IL-1 antagonist; TMP; TPO mimetic peptide; EPO mimetic peptide; EMP; VEGF antagonist; MMP inhibitor; antiinflammatory; antitumour; immunosuppressive; cytostatic; antiinflammatory; antitutus; interleuketic; ophthalmological; antiinaemic; anorectic; antiinfertility; haemostatic; dermatological;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuroprotective; inflammatory disease; autoimmune disease; tumour growth; cancer; rheumatoid arthritis; diabetic retinopathy; infertility; obesity; sleep discorder; neurological degenerative disease; anaemia; thrombocytopaenia; metastatic tumour; systemic lupus erythematosus;
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abb73178
ABB73177 Length: 15 October 13, 2004 13:39 Found using 'claim36' (zara371.key)
        The present invention describes a vehicle-peptide molecule (I) or its cmultimers. (I) can have antianflammatory, antitumour, immunosuppressive, cytostatic, antirheumatic, antianthritic, antidiabetic, ophthalmological, antianaemic, anorectic, antianfertility, haemostatic, dermatological and consuppressive, and the present of the present of the present of the proposes. (I) is useful for composing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of their constant of diagnostic kit to detect the presence of their corrections of interest in a biological sample. Additionally, (I) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders, confinition compounds are useful for treating disorders characterised by low red blood cell levels such as anaemia. The TPO-mimetic composition
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abb73178; Mdm/hdm antagonist peptide SEQ ID NO:139.
(from "claim36agg.pep")
TOIG of: abb73178 check: 9428 from: 1 to: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; antirheumatic; antiarthritic; antidiabetic; ophthalmological; antianaemic; ancrectic; attiinfertility; haemostatic; dermatological; neuroprotective; inflammatory disease; autoimmune disease; tumour growth; cancer; rheumatoid arthritis; diabetic retinopathy; infertility; obesity; sleep disorder; neurological degenerative disease; anaemia; thrombocytopaenia; metastatic tumour; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                              Claim 39; Page 53; 176pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                       Novel vehicle-peptide molecule or its multimers useful for treating inflammatory and autoimmune diseases, cancer, rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-MAY-2000; 2000US-00563286.
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abb73179; Mdm/hdm antagonist peptide SEQ ID NO:140 (from "claim36ags.pep")
TOIG of: abb73179 check: 8833 from: 1 to: 15
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neuroprotective activities. (I) can be used as a therapeut prophylactic agent as well as for screening purposes. (I) diagnosing diseases characterised by dysfunction of their
                                                  multimers. (I) can have antiinflammatory, antitumour, immunosuppressicytostatic, antirheumatic, antiarthritic, antidiabetic, ophthalmologiantianaemic, anorectic, antiinfertility, haemostatic, dermatological
                                                                                                                                                                                  Novel vehicle-peptide molecule or its multimers useful for treating inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility.
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                                                                                                             The present invention describes a vehicle-peptide molecule (I) or its
                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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ng 'claim36'
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                                                                                                                                                                                                                                                                                                                                             AMGEN INC
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                     (I) can be used as a therapeutic or se for screening purposes. (I) is us
                                                                                                                                                                                                                                                                                                       Boone TC,
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                                                                                              immunosuppressive,
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l match found in sequence:
abb92234 ; Herbicidally active polypeptide SEQ ID NO
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TOIG of: abb92234 check: 9732 from: 1 to: 440
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                                              amino acid sequences from non-plant organisms using suitable search parameters, where plant sequences having an E-value greater by a factor of 3 than the E-value of most similar non-plant sequences are selected. The polypeptides or nucleic acids encoding them are useful for
                                                                                                                                                           The invention relates to identifying target proteins (ABB90790-ABB94016) for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or
                                                                                                                                                                                                                                                                                                                                                      Identifying plant target proteins for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or amino acid sequences from non-plant
                                                                                                                                                                                                                                                                           Claim
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abb92409; Herbicidally active polypeptide SEQ ID
(from "claim36ags.pep")
TOIG of: abb92409 check: 2170 from: 1 to: 476
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             ABB92409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Found using
                                                                                  The invention relates to identifying target proteins (ABB90790-ABB94016) for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or amino acid sequences from non-plant organisms using suitable search parameters, where plant sequences having an E-value greater by a factor of 3 than the E-value of most similar non-plant sequences are selected. The polypeptides or nucleic acids encoding them are useful for identifying modulators. The identified modulators are useful as
                                                                                                                                                                                                                                                                   Identifying plant target proteins for herbicidally active compounds, comprising aligning and comparing nucleic acid or amino acid sequences from plant with nucleic acid or amino acid sequences from non-plant
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                                           Sequence 476
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Arabidopsis thaliana.
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using
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KTEYKESLPAPENNDLYKGKIWCVGNNTCDPIQRGGPCQKPDLTVLHASYAPESYWAQFR
400 408
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Length: 476
ng 'claim36'
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446
adb61841 ; Peptide Seq ID81 (from "claim36ags.pep")
                                    match found in sequence
                                                                                                NQSSQCYFNGLARETTTNPGNERCKFPSVTL
                        related to inhibitors
                    of apoptosis
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TOIG of: adb61841
                                                                                      04-DEC-2003
                                                                                           ADB61841 standard;
                                                                                        ADB61841;
                                                                                      (first entry)
                                                                                              check: 3739
                                                                                           peptide;
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autoimmune disorder; neurodegenerative disease; apoptotic response; systemic lupus erythematosis; multiple sclerosis; viral infection; herpes virus; poxvirus; adenovirus; inhibitor of apoptosis; IAP; XII HIAP1; cIAP2; CIAP1; RING zinc finger; caspase-3; caspase-7; caspase-9; cytostatic; neoplasm; leukaemia; colon carcinoma; cervical cancer; uterine cancer; testicular cancer; small cell lung carcinoma; uterine cancer; renal cell carcinoma; Wilm's tumour apoptosis pathway; embryonic development; viral pathogenesis; cancer; baculovirus inhibitor of apoptosis repeat domain; BIR domain; Peptide Seq ID81 related to inhibitors of apoptosis.

09-NOV-2001; 2001US-0332300P. 08-APR-2002; 2002US-0370934P. Unidentified Boudreault A, 12-NOV-2002; 2002WO-CA001738 WO2003040172-A2 (AEGE-) AEGERA THERAPEUTICS INC 15-MAY-2003 Korneluk RG, Ľа Casse 'n, Liston

cancer and other neoplasms Polypeptide capable of forming a complex with a polypeptide comprising baculovirus inhibitor of apoptosis repeat domain useful for treating WPI; 2003-513532/48.

Disclosure; Page 13; 53pp; English.

(EIR) Journal The application pathway is known to play a critical role in cembryonic development, viral pathway is known to play a critical role in cembryonic development, viral pathway is known to play a critical role in cembryonic development of diseases. The failure of the apoptotic response has been implicated in the development of cancer, autoimmune disorders (for example systemic lupus erythematosis and multiple sclerosis) and viral infections (including herpes virus, poxvirus and adenovirus. The critical control of proteins possessing one or more baculovirus IAP repeat (BIR) domains. Human IAPs, XIAP, HIAP1 (CIAP1) and HIAP2 (CIAP1) all posses three BIR domains and carboxy terminal RING zinc fingers. The IAPs bind and inhibit caspases -3, -7 and -9 which are proteases involved in the initiation of apoptosis. Compounds which inhibit the activity of IAPs may therefore have cytostatic activity through the enhancement of apoptosis. The polypeptides of the invention care candidate paptide ligands for binding to the BIR domain of IAPs. They may be useful for the treatment of cancer and other neoplasms, such as leukaemias, colon carcinoma, cervical cancer, terrine cancer, testicular cancer, small cell lung carcinoma, uterine cancer, renal carcinoma and wilm's tumour, and for enhancing apoptosis. The present sequence is that of a peptide which is suggested as a possible peptide for fusing to This invention relates to a substantially pure polypeptide ength of less than 100 amino acids and capable of forming a complex with polypeptide that includes a baculovirus inhibitor of apoptosis repeat

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VWCVAVDGADEAELGQALNFACGRSNATCAALAPGGECYAPVTVTWHASYAPSSYWAQFR
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